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1: /cgn2_6/ptcdata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptcdata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptcdata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptcdata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptcdata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptcdata/1/ina/backfiles1.seq:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Sequence 17468, A
Sequence 13589, A
Sequence 17009, A
Sequence 11808, A
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ALIGNMENTS

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US-09-949-016-17468/c
US-09-949-016-17468/c
Sequence 17468, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICATT: VENTER, J. Craig et al.
APPLICATT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISBASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
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; ORGANISM: Human
US-09-949-016-203106
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PRIOR PILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FeetSEQ for Windows Version 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 203106, Application US/09949016
PATENT NO. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
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SEQ ID NO 203106
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Best Local Similarity
Matches 41; Conserv
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CURRENT FILING DATE: 2000-04-14
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OF DETECTION AND USES THEREOF
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT ETLING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILLING DATE: 2000-10-03
PRIOR FILLING DATE: 2000-10-03
PRIOR FILLING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILLING DATE: 2000-09-08
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US-09-513-999C-13589
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION: APPLICANT: VENTER, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17009, Application US/09949016 Patent No. 6812339
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LENGTH: 182
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CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
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SOFTWARE: Patent.pm
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NUMBER OF SEQ ID NOS: 207012
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FEATURE:
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(88906)
OTHER INFORMATION: n = A,T,C or
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85.4%;
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Pred. No. 0.0012;
0; Mismatches !
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Pred. No. 0.0022;
0; Mismatches
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OF DETECTI
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GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-30

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498
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US-09-949-016-17009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FRASTSEQ for Windows Version 4.0
SEQ ID NO 11808
LENGTH: 636591
                                                                                                                                                                                                                                                                                                                                                       Sequence 13388, Application US/09949016 Patent No. 6812339
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LENGTH: 205163
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241.755
PRIOR APPLICATION NUMBER: 60/241.755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
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Best Local (
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13388
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NAME/KEY: misc_feature
LOCATION: (1)...(636591)
OTHER INFORMATION: n = A,T,C
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Similarity 80.4%;
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Pred. No. 0.0079;
0; Mismatches 1;
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Pred. No. 0.014;
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CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEO ID NOS: 207012
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                                                                             PATENT OF 16812339

PATENT NO. 6812339

PATENT NO. 6812339

GENERAL IMPORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                               Sequence 22414, Application US/09949016
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LENGTH: 601
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SEQ ID NO 22414
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                 PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: PASTSEQ for Windows Version 4.0
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LOCATION: (1)...(636591)
OTHER INFORMATION: n = A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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Local Similarity 80.4%;
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Pred. No. 0.0041;
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Pred. No. 0.014;
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; ORGANISM: Human US-09-949-016-154219
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                                                                                           NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 154219
LENGTH: 601
                                                                                                                                                                                                                                                                                                                                                                                                      Patent No. 6812339 GENERAL INFORMATION:
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SOFTWARE: FastSEQ for
SEQ ID NO 154217
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Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 154219, Application US/09949016
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Best Local :
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                                                                                                                                                                 APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILLING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILLING DATE: 2000-10-20
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PRIOR APPLICATION NUMBER: 60/231,498
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TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Human
                                                                           TYPE: DNA
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39; Conserv
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Pred. No. 0.0041;
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Pred. No. 0.0041;
0; Mismatches
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Length 601;
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84.8%;

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43449 ACCTCTTTATACACACACACACACACACACACACGAAATGTATG 43494

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PRIOR APPLICATION NUMBER: 60/241.755
PRIOR FILING DATE: 2000-10-20
PRIOR PILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR PILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.
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US-09-949-016-11957
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16064
LENGTH: 70131
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
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Patent No. 6812339
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                                                               Matches
                                                                                                  Query Match
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Best Local
                                                                                                                                         ORGANISM: Human
-09-949-016-16064
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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CURRENT FILING DATE: 2000-04-14
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ORGANISM: Human
                                                                                                                                                                                  TYPE: DNA
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39; Conserv
4 ATCTCTTTCTTTACACACACACACACACACACACACAAAATATCTG 49
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nilarity 84.8%;
Conservative
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                                                           Score 34.8; DB Pred. No. 0.01; 0; Mismatches
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0; Mismatches 7;
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GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT FAPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-19-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-19-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-33,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-08
PRIOR FILING DATE: 207012
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 142944
LENGTH: 601
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                                                                                                                                       Matches
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ORGANISM: Human
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37;
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37; Conserv
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90.2%;
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OF DETECTION
                                                                                                                                                                                                        Length 601;
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RESULT 15 US-09-949-016-142945

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Database
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Maximum Match 100%
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Maximum DB
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                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Gapop 10.0 , Gapext 1.0
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19:
20:
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/ Ggn2_6/ptodata/1/pubpna/US09B_PUBCOMB.seq:*
/ Ggn2_6/ptodata/1/pubpna/US09B_PUBCOMB.seq:*
/ Ggn2_6/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
/ Ggn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
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/cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/USO6_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/USO6_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/USO7_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/USO8_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/USO8_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/USO8_PUBCOMB.seq:*
SUMMARIES
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100.0 2085 79.2 10619 79.2 10619 79.2 10619 79.2 10619 79.2 10619 75.7 400 73.3 154	Query Match
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Sequence 5, Appli Sequence 2, Appli Sequence 44, Appli Sequence 2, Appli Sequence 2, Appli Sequence 3039, Ap Sequence 10920, A	Description

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-09-956-712-1 -10-633-913-1	-10-322-281-66	-027-632-180	-10-027-632-18056	-10-674-124A-1721	-10-674-124A-39	-10-674-124A-5849	-10-674-124A-1052	0-674-124A-19	-10-674-124A-10	-10-674-12	-10-674-124A-94	-10-715-066-	-10-027-632-14946	-10-027-632-14946	-10-027-632-14	-10-027-632-14946	-10-027-632-149	-10-027-632-14945	-10-027-632-12816	-10-027-632-12816	-10-674-124A-2524	-10-674-124A-2322	-10-674-124A-2525	-10-674-124A-1202	-10-027-632-28241	-10-027-632-28	-10-087-192-496	-10-674-124A-12	-10-756-149-116	-10-027-632-25946	-10-027-632-2594	-10-674-124A-1821	US-10-717-597-322	US-10-655-847-4	0-160-	S-10-357-
Sequence 10, Appl Sequence 10, Appl	667,	180561	18056	17215,	e 3935,	e 5849,	e 10528,	e 19062,	e 10989,	e 6272	e 9430,	е 3, Арр					Sequence 149460,		_		25249,	23225,	25250,	Sequence 12025, A	282410	e 28241	e 496, A	æ	e 1160,	259460	σ	18213,	e 322	O	e 4, Appl	e 55243,

ALIGNMENTS

US-09-909-317-5

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; Sequence 5, Application US/09909317
publication No. US20040152075A1
GENERAL INFORMATION:
APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Decreme I. Rotter (Inventor)
APPLICANT: Decreme I. Rotter (Inventor)
APPLICANT: Decreme I. Rotter (Inventor)
APPLICANT: Data: 18810-82152
CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR APPLICATION NUMBER: 09/280,317
PRIOR APPLICATION NUMBER: 09/280,31
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; FEATURE:
; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
US-10-311-455-44
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US-10-311-455-44/c
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US-10-239-676-2/c
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NUMBER OF SEQ ID NOS: 2
SEQ ID NO 2
LENGTH: 10619
TYPE: DNA
                                                                                                                                                                     SEQ ID NO 44
LENGTH: 10619
TYPE: DNA
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Publication No. US20030082609A1
    Matches
                 Query Match
Best Local Similarity
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Best Local
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                                                                                                                                                                                                                          APPLICANT: OLEK, Alexander
APPLICANT: DIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Determ
TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Determ
TITLE OF INVENTION: Cytosine methylation
FILE REFERENCE: 5013.1014
CURRENT APPLICATION NUMBER: US/10/311,455
CURRENT FILING DATE: 2001-07-02
PRIOR APPLICATION NUMBER: DCT/EP01/07537
PRIOR PILING DATE: 2001-07-02
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
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APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with Gene Regulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DE 10043826.1
PRIOR FILING DATE: 2001-04-06
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DE 10019058.8
DE 10019173.8
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CURRENT APPLICATION NUMBER: US/10/239,676
CURRENT FILING DATE: 2002-09-24
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                                                                                                                                                ORGANISM: Artificial Sequence
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2000-04-07
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Similarity 97.6%;
    Conservative
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                    79.2%;
Score 40.4; DB 19
Pred. No. 0.00038;
0; Mismatches 1
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Pred. No. 0.00038;
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                                          DB 15;
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    Indels
                                        Length 10619;
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GENERAL INFORMATION:

APPLICANT: OLEK, Alexander

APPLICANT: PIEPENBROCK, Christian

APPLICANT: BERLIN, Kurt

ITILE OF INVENTION: Diagnosis of Diseases Associated with

ITILE OF INVENTION: Diagnosis of Diseases Associated with

ITILE REFERENCE: 5013.1008

CURRENT APPLICATION NUMBER: US/10/240,589C

CURRENT FILING DATE: 2003-09-02

PRIOR APPLICATION NUMBER: CT/FE01/03972

PRIOR APPLICATION NUMBER: DE 10019058 8

PRIOR FILING DATE: 2000-04-06

PRIOR FILING DATE: 2000-04-07

PRIOR APPLICATION NUMBER: DE 10019173.8

PRIOR APPLICATION NUMBER: DE 10032529.7

PRIOR APPLICATION NUMBER: DE 10032529.7

PRIOR APPLICATION NUMBER: DE 10043826.1
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US-10-240-453-2/c
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CURRENT FILING DATE: 2002-10-02
PRIOR APPLICATION NUMBER: PCT/EP01/03973
PRIOR FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR PILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
PRIOR FILING DATE: 2000-06-30
  SEQ
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Best Local S
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PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 148
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA TITLE OF INVENTION: Transcription TITLE OF INVENTION: by Means of Assessing the Methylation Status of Genes Associated TITLE OF INVENTION: with DNA TRanscription FILE REFERENCE: 5013.1009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
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Local Similarity 97.6%;
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Pred. No. 0.00038;
0; Mismatches 1;
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RESULT 7
US-10-674-124A-10920
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CURRENT APPLICATION NUMBER: US/10/674,124A
CURRENT FILING DATE 2003-09-26
PRIOR APPLICATION NUMBER: 10/257.511
PRIOR REPLICATION NUMBER: 10/257.511
PRIOR REPLICATION NUMBER: PCT/JP00/07621
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2000-10-30
PRIOR APPLICATION NUMBER: JP2000-112699
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR FILING DATE: 2002-99-28
PRIOR FILING DATE: 2002-99-28
PRIOR FILING DATE: 2002-99-28
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Matches
                                       Sequence 10920, Application US/10674124A Publication No. US20040197797A1 GENERAL INFORMATION:
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LENGTH: 400
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APPLICANT: INOKO, H: APPLICANT: TAMIYA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 27110
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PRIOR FILING DATE: 2002-12-09
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APPLICANT: TAMIYA, Gen
TITLE OF INVENTION: GENE MAPPING METHOD USING MICROSATELLITE
TITLE OF INVENTION: GENETIC POLYMORPHISM MARKERS
                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Distance between 3'-terminus of neighbour sequence OTHER INFORMATION: sequence listing upward to telomere on chrosomal stother INFORMATION: 5'-terminus of this base sequence; 18142
                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
OTHER INFORMATION: Distance between a terminus base of telomere
OTHER INFORMATION: chromosomal short arm and 5'-terminus of this
OTHER INFORMATION: sequence: 84667136
                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Located on chromosome 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: AC010744.4_97739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 10619
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                     TCTCTCTCTATACACACACACACACACACACACACAATATATATG
                                                                                                                                                                                                                          TCTCTTTCTTTACACACACACACACACACACACACAAAATATCTG 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  75.7%;
llarity 91.1%;
Conservative
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                 Hidetoshi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    79.2%;
97.6%;
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Pred. No. 0.00
0; Mismatches
                                                                                                                                                                                                                                                                 Score 38.6; DB 19; Length 400; Pred. No. 0.00087; 0; Mismatches 4; Indels 0
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s base
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US-10-357-930-55243
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PRIOR APPLICATION NUMBER: 10/257,511
PRIOR TILING DATE: 2003-03-07
PRIOR PPLICATION NUMBER: PCT/JP00/07621
PRIOR FILING DATE: 2000-10-30
PRIOR APPLICATION NUMBER: JP2000-112699
PRIOR FILING DATE: 2000-04-13
PRIOR PILING DATE: 2000-04-13
PRIOR PILING DATE: 2002-09-28
PRIOR PILING DATE: 2002-09-28
PRIOR PILING DATE: 2002-09-28
PRIOR PILING DATE: 2002-12-09
PRIOR FILING DATE: 2002-12-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 55243, Application US/10357930 Publication No. US20040259086A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 27110
SEQ ID NO 10920
LENGTH: 154
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Schlegel. Robert
APPLICANT: Endege, Wilson
APPLICANT: Monahan, John
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS,
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREV
FILE REFERENCE: MRI-007BCN
FILE REFERENCE: MRI-007BC
                                                                                                                                                                                                                                                           PRIOR
                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/189,862
PRIOR FILING DATE: 2000-03-16
PRIOR APPLICATION NUMBER: 60/207,454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 09/785,276 PRIOR FILING DATE: 2003-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/357,930 CURRENT FILING DATE: 2003-02-04
                                        NUMBER
                                                                            PRIOR APPLICATION NUMBER: 60/255,281
PRIOR FILING DATE: 2000-12-13
                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/219,007 PRIOR FILING DATE: 2000-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/183,319
PRIOR FILING DATE: 2000-02-17
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                                                                                                                                                                                                                                                                                                                                                PRIOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Distance between 3'-terminus of neighbour sequence of OTHER INFORMATION: sequence listing upward to telomere on chrosomal short arm OTHER INFORMATION: 5'-terminus of this base sequence: 107348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Distance between a terminus base of telomere on OTHER INFORMATION: chromosomal short arm and 5'-terminus of this b OTHER INFORMATION: sequence : 131608898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Located on chromosome FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
OTHER INFORMATION: D6S988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
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                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/211,314 FILING DATE: 2000-06-09
                                                                                                                                                                                                                                                                                                                                                FILING DATE: 2000-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             76 TCTCTTTCCTGACACACACACACACACACACACACACACTCTCTCAT 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 TCTCTTTCTTTACACACACACACACACACACACAAAATATCTGAT
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                                            SEQ ID NOS: 62232
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FastSEQ
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for Windows Version 4.0
                                                                                                                                                                                                                                                      2000-06-09
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Pred. No. 0.0019;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ASSESSMENT, PREVENTION, AND
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SEQ ID NO 55243

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                                                                 US-10-655-847-4
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                                                                                                                                                                                                                   Sequence 4, Application US/10655847
Publication No. US20040063129A1
GENERAL INFORMATION:
APPLICANT: William Gaarde
APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
CURRENT APPLICATION NUMBER: US/10/655,847
CURRENT FILING DATE: 2003-09-05
CURRENT FILING DATE: 2003-09-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
Query Match
Best Local Similarity
Matches 41; Conserv
                                                                                                                                                     NUMBER OF SEQ ID NOS: 296
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/10160807 Publication No. US20030224514A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION FILE REFERENCE: RTS-0189 CURRENT PROLICATION NUMBER: US/10/160,807 CURRENT FILING DATE: 2002-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-160-807-4
                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US/10/160,807 PRIOR FILING DATE: 2003-09-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: 31
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                     LENGTH: 104245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 104245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 / Match 72.2%;
Local Similarity 85.4%;
nes 41; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 72.5%;
Local Similarity 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 161 TCTCTCTCTACACACACACACACACACACACACAAATTAGCTG 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 40;
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 Conservative
               72.2%;
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Pred. No. 0.0034;
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Score 36.8; DB 18; Length 104245; Pred. No. 0.01; 0; Mismatches 7; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 36.8; DB Pred. No. 0.01; O; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 104245;
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Gaps
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; SOFTWARE: PatentIn version 3.2; SEQ ID NO 322; LENGTH: 170245; TYPE: DNA; ORGANISM: Homo sapiens US-10-717-597-322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18213, Application US/10674124A Publication No. US20040197797A1 GENERAL INFORMATION:
                                                                                                  SEQ ID NO 18213
LENGTH: 419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 72.2%;
Best Local Similarity 85.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 322, Appublication No.
                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: GENE MAPPING METHOD USING MICROSATELLITE TITLE OF INVENTION: GENETIC POLYMORPHISM MARKERS FILE REFERENCE: ORIN-003CIP CURRENT APPLICATION NUMBER: US/10/674,124A CURRENT FILING DATE: 2003-09-26 PRIOR APPLICATION NUMBER: 10/257,511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: AM101080L
CURRENT APPLICATION NUMBER: US/10/717,597
CURRENT FILING DATE: 2003-11-21
                                                                                                                                        PRIOR APPLICATION NUMBER: JP2002-383869
PRIOR FILING DATE: 2002-12-09
NUMBER OF SEQ ID NOS: 27110
                                                                                                                                                                                                  PRIOR FILING DATE: 2003-03-07
PRIOR APPLICATION NUMBER: PCT/JP00/07621
PRIOR FILING DATE: 2000-10-30
PRIOR APPLICATION NUMBER: JP2000-112699
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR APPLICATION NUMBER: JP2002-327516
PRIOR FILING DATE: 2002-09-28
PRIOR FILING DATE: 2002-09-28
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NUMBER OF SEQ ID NOS: 4904
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PRIOR FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: US 60/427,982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Stover, Jennifer A. TITLE OF INVENTION: METHODS FOR DIAGNOSING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: INOKO, Hidetoshi
APPLICANT: TAMIYA, Gen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Wyeth
FEATURE:
OTHER INFORMATION: AC004802.1_36841
FEATURE:
                                                       TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
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Trepicchio, William L.
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Twine, Natalie
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o. US20040110221A1
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Pred. No. 0.011;
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RESULT 14
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US-10-027-632-259460/c
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
      Sequence 259460, Application US/10027632

Publication No. US20030204075A9

GENERAL INFORMATION:

APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide

TITLE OF INVENTION: Polymorphisms in the Human Genome

FILE REFERENCE: 108827.129

CURRENT APPLICATION NUMBER: US/10/027,632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 259460
LENGTH: 657
TYPE: DNA
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Distance between 3'-terminus of neighbour sequence of OTHER INFORMATION: sequence listing upward to telomere on chrosomal short arm and OTHER INFORMATION: 5'-terminus of this base sequence : 223511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Distance between a terminus base of telomere on OTHER INFORMATION: chromosomal short arm and 5'-terminus of this base OTHER INFORMATION: sequence : 1189010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE
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83.7%;
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                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/756,149
CURRENT FILING DATE: 2004-01-12
NUMBER OF SEQ ID NOS: 5818
SOPTWARE: Patentin version 3.2
SEQ ID NO 1160
LENGTH: 754
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1160, Application US/10756149
Publication No. US20050181375A1
GENERAL INFORMATION:
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                                                                      Matches
                                                                                                     Query Match
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Best Local Similarity
                                                                                                                                                                                                                                                                                                              APPLICANT: Aziz, Natasha
APPLICANT: Zlotnik, Albert
TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSIS OF METASTATIC CANCER, COMPOSITIONS AND
TITLE OF INVENTION: METHODS OF SCREENING FOR MODULATORS OF METASTATIC CANCER
                                                                                                                                                                                                                                                                                                     FILE REFERENCE: file
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                                                                                                                                                         ORGANISM: Homo Sapiens
                                                                                                                                                                           TYPE: DNA
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                                                                                    Local
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APPLICATION NUMBER: US 60/146,002
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311
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                       5 TCTCTTTCTTTACACACACACACACACACACACAAAATATCT 48
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100.0%; Pred. No.
tive 0; Mismatch
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1: /cgn2 6/ptodata/1
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length: 2000000000
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US-09-280-181B-1
US-09-248-796A-1081B
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US-09-949-016-111397
US-09-949-016-12809
US-09-949-016-13408
US-09-949-016-13408
US-09-949-016-137497
US-09-949-016-15508
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Sequence 1, Appli
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Sequence 17107, A
Sequence 13548, A
Sequence 13548, A
Sequence 14693, A
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Sequence 14010, A
Sequence 10881, A
Sequence 80901, A
Sequence 111397,
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US-09-280-181B-1
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Minimum Maximum

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Scoring table: Sequence: Title: Perfect score:

Searched:

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PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14010
                                                                                                                                                                                                                                                                           Sequence 14010, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
CURRENT FILING DATE: 2000-04-14
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CURRENT FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 22
TYPE: DNA
ORGANISM: Homo sapiens
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Patent NO. 6280941
GENERAL INFORMATION:
APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
TITLE OF INVENTION: Genetic Marker Test for Lupus
FILE REFERENCE: P07 41735
LENGTH: 34408
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOPTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 80901
LENGTH: 601
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; LOCATION: (1) ... (34408)
; OTHER INFORMATION: n = A,T,
US-09-949-016-14010
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US-09-248-796A-10881
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US-09-949-016-80901
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; ORGANISM: Candida albicans
US-09-248-796A-10881
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES
FILE REFERENCE: CL001307
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TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 10881
LENGTH: 366
TYDES: NUMBER: NUMBER: US 60/096,709
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Best Local Similarity 95.0%;
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                    ORGANISM: Human
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Similarity 90.5%;
19; Conservative
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Pred. No. 1.
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Pred. No. 1e+02;
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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APPLICANT: Keith Weinstock et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.132
CURRENT PELICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-08-13
NUMBER: OF SEQ ID NOS: 28208
                                                                    Query Match
Best Local S
Matches 19
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LENGTH: 711
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 111397
LENGTH: 601
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Best Local Similarity
Matches 19; Conserv
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                          TYPE: DNA ORGANISM: Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Human
                                                                                         Local Similarity
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                                                                    Score 17.8; DB 4;
Pred. No. 1.2e+02;
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Pred. No. 1.2e+02
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pred. No. 1.2e+02;
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                                                                  1.2e+02;
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RESULT 9
US-09-949-016-14083/c
; Sequence 14083, Application US/09949016
; Patent No. 6812339
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; ORGANISM: Human
US-09-949-016-14816
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US-09-949-016-14816
                                                                                                                                                                                                                                                                                                            ; ORGANISM: Human
US-09-949-016-13408
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                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 13408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13408, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/237,768 PRIOR FILING DATE: 2000-10-03
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR PPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                   LENGTH: 49378
TYPE: DNA
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                                                                                                                                                                                                                 80.9%;
1 Similarity 90.5%;
19; Conservation
                                                                                                                                                           2 ATTCCCCATCTCTCTTTCTTT 22
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Pred. No. 1.9e+02;
0; Mismatches 2;
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OF DETECTION AND USES THEREOF
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Query Match
Best Local Similarity
Watches 18; Conserva
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US-09-949-016-69074/c
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CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/231,755

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                   Sequence 69074, Application US/09949016

Patent No. 6812339

GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 69074
LENGTH: 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FRATSEQ for Windows Version 4.0
SEQ ID NO 14083
LENGTH: 199471
TYPE: DNA
ORGANISM: Human
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APPLICANT: VENTER, J.
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PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: (1)...(199471)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                                                                        TYPE: DNA
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331 ATTCCCCATCTCTCTTCCT 313
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1 Similarity 90.5%;
19; Conservative
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                         ATTCCCCATCTCTCTTTCT 20
                                                                                     Conservative
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                                                                                                        79.1%;
94.7%;
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                                                                                                        Score 17.4; DB 4;
Pred. No. 1.8e+02;
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Pred. No. 2.3e+02;
                                                                                     Mismatches
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RESULT 11 US-09-949-016-137497/c ; Sequence 137497, Application US/09949016 ; Patent No. 6812339.

GENERAL INFORMATION:

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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR RILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 137497
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                Sequence 12500, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
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Patent No. 6812339
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Best Local :
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APPLICANT: VENTER, J.
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PILING DATE: 2000-10-03
PRIOR PPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
CURRENT APPLICATION NUMBER: US/09/949,016
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Human
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94.7%;
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94.7%;
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Pred. No. 2.6e
0; Mismatches
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Pred. No. 1.8e+02;
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; PRIOR FILING DATE: 2000-09-08; NUMBER OF SEQ ID NOS: 207012; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 16092; LENGTH: 53337
; TYPE: DNA; ORGANISM: Human
US-09-949-016-16092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: Fast SEQ for Windows Version
; SEQ ID NO 12500
; LENGTH: 53336
; TYPE: DNA
; ORGANISM: Human
US-09-949-016-12500
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Best Local S
Matches 18
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Patent No. 6812339
GENERAL INFORMATION:
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Patent No. 6812339
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
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PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/231,498
                                                                                                                                                                                                                                                                                                                                                                  8772
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18; Conserv
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94.7%;
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Pred. No. 2.9e+02;
0; Mismatches 1;
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Pred. No. 2.9e+02;
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OF DETECTION AND USES THEREOF
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Maximum DB seq length: 2000000000
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Gapop 10.0 , Gapext 1.0
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22
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1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
3: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
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/ Ggn2_6/ptodata/2/pubpna/USO7_NEW_PUB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO8_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO8_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO8_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO9A_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO9A_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO9C_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO9C_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO9C_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USO9C_PUBCOMB.seq:*
/ Cgn2_6/ptodata/2/pubpna/USOOC_PUBCOMB.seq:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

c 7 19	c 6 19	ຫ	C 4 19	c 3 20		22	Result No. Score
86.4	86.4	86.4	86.4	90.9	90.9	0	Query
_	32221	32189	32189	650	650		Query Match Length DB ID
14	9	14				11	. BB
14 US-10-079-854-377	US-09-764-878-377	US-10-079-854-379	US-09-764-878-379	US-10-027-632-190184	US-10-027-632-190184	US-09-909-317-1	DB ID
Sequence 377. App	Sequence 377, App	Sequence 379, App	Sequence 379, App	Sequence 190184,	Sequence 190184,	Sequence 1, Appli	Description

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10-767-701- 10-203-319 <i>1</i> 10-203-319 <i>1</i>	1 ()	L0-027-632-17476)-741-600-26828	-027-632	0-415-05)-719-993-67)-459-262A-3	10-240-485-15	10-027-632-20691 10-311-455-1925	10-425-115	10-027-632-	10-027-632	-10-027-632	-10-027-632-	US-10-424-599-31004 US-10-674-124A-23460	-10-741-601-561	-10-085-783A-2643	-10-242-535A	-10-741-600-176	566 66	-10-741-600-	10-433-793-11	-10-221-613-19	-10-311-455-	-10-424-599-5903	-10-424-599-92	-10-183-687-2	US-10-180-375-7
Sequence 12634, A Sequence 22, Appl Sequence 24, Appl	Sequence 101998, Sequence 101998, Sequence 94947, A	Sequ	Sequence 174763,	Sequence 5, Appli	Sequence 6760, Ap Sequence 3, Appli	O I	e 206	Sequence 206914,	206915,		"	206913	Seguence 31004, A	5616,	26433,	Sequence 26433, A	17685,	Sequence 5660, Ap	Sequence 17886, A	e 11	19	e 14:	59	921	equence 23, A	Sequence 7, Appli

ALIGNMENTS

RESULT 1 US-09-909-317-1

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                                                                                                                                                      ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-909-317-1
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                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 22
                                                                           Best Loc
Matches
                                                                                                Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                      APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
IITLE OF INVENTION: Genetic Marker Test for Lupus
FILE REFERENCE: 18810-82152
CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
CURRENT FILING DATE: 2001-07-18
                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 09/280,181 PRIOR FILING DATE: 1999-03-29
                               1 GATTCCCCATCTCTCTTTT 22
                                                                             22;
                                                                                                Similarity
                                                                           100.0%; Score 22; DB ilarity 100.0%; Pred. No. 6.7; Conservative 0; Mismatches
22
                                                                                                                   DB 11; Length 22;
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US-10-027-632-190184/c
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US-10-027-632-190184/c
; SEQ ID NO 190184
; LENGTH: 650
TYPE: DNA
                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
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Publication No. US20020198371A1
GENERAL INFORMATION:
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LENGTH: 650
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Best Local (
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
                                                                        PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4
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PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/218,006 PRIOR FILING DATE: 2000-07-12
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PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
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PRIOR FILING DATE: 2000-07-12
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CURRENT FILING DATE: 2002-04-30
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                                                                                                                                                                      APPLICATION NUMBER: US 60/156,358 FILING DATE: 1999-09-28
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Pred. No.
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56;
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WS-09-764-878-377/c
US-09-764-878-377/c
; Sequence 377, Application US/09764878
; Patent No. US20020090615A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; SOFTWARE: PatentIn Ver. ; SEQ ID NO 379 ; LENGTH: 32189 ; TYPE: DNA CORGANISM: Homo sapiens
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US-09-764-878-379/c
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                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 379, Application US/10079854 Publication No. US20030054368A1 GENERAL INFORMATION:
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA121
CURRENT APPLICATION NUMBER: US/09/764,878
CURRENT FILING DATE: 2001-01-17
                                                                                                                                                                                                                                                                                                           Prior Application removed - See File Wrapper NUMBER OF SEQ ID NOS: 428
                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/079,854
CURRENT FILING DATE: 2002-02-22
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA121C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 428
                                                                                                                                                                                                                                                   LENGTH: 32189
TYPE: DNA
                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
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                                                                                                      19817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19817
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19; Conserv
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                                                                                                        rccccarcrcrcrrrcrrr 19799
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100.0%; Pred. No. .
''ve 0; Mismatches
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100.0%; Pre
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100.0%; Pr.
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0; Mismatches
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Pred. No.
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hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                      and Antibodies
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Sequence 7, Application US/10180375
Publication No. US20030126638A1
GENERAL INFORMATION:
APPLICANT: Allen, William B.
APPLICANT: Cahoon, Rebecca E.
APPLICANT: Famodu, Omolayo O.
APPLICANT: Harvell, Leslie T.
APPLICANT: Helentjaris, Timothy
APPLICANT: Li, Changjiang
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, NAME/KEY: SITE
; LOCATION: (7464)
; OTHER INFORMATION: n equals a,t,g, or
US-09-764-878-377
                                                                                                                                                                                             US-10-180-375-7/c
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Best Local S
Matches 19
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PAL21C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 377
LENGTH: 32221
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SEQ ID NO 377
LENGTH: 32221
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            APPLICANT:
APPLICANT:
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CURRENT FILING DATE: 2002-02-22
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA121
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CURRENT FILING DATE: 2001-01-17
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NAME/KEY: misc_feature
LOCATION: (7464)
OTHER INFORMATION: n equals a,t,g, or
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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les 19; Conservative
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100.0%; Pr
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100.0%; Pr
            Igor Cunha
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+(
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hes 0;
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US-10-424-599-92893/c

Sequence 92893, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:

APPLICANT: La Rosa Thomas APPLICANT: Kovalic David APPLICANT: Zhou Yihua

APPLICANT:

Cao Yongwei

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US-10-183-687-23
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SEQ ID NO 23
LENGTH: 1334
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TITLE OF INVENTION: Alteration Of Oil Traits In Plants
FILE REFERENCE: BB1458 US NA1
CURRENT APPLICATION NUMBER: US/10/180,375
CURRENT FILING DATE: 2002-06-26
NUMBER OF SEQ ID NOS: 222
SOFTWARE: Microsoft Office 97
SEQ ID NO 7
LENGTH: 1334
                                                       Matches
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                                                                     Query Match
Best Local Similarity
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FILE REFERENCE: BB1458 US NA
CURRENT APPLICATION NUMBER: US/10/183,687
CURRENT FILING DATE: 2002-06-27
PRIOR FILING DATE: 2002-06-29
NUMBER OF SEO ID NOC. 572
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Best Local Similarity
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APPLICANT:
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                                                                                                                                               TYPE: DNA
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1238 GATTCCCAATCTCTCTGTCTTT 1217
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                  1 GATTCCCCATCTCTCTTTCTTT 22
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Shen, Bo
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Harvell, Leslie T.
Jones, Todd
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Oliveira, Igor Cunha
Sakai, Hajime
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Klein, Ted
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                                                         Conservative
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90.9%;
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90.9%; Pred. No. 1
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                                                                     Score 18.8;
Pred. No. 1.
                                                         Mismatches
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Sequence 1419, Application US/10311455
Publication No. US20030143606A1
GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: PIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated wit
TITLE OF INVENTION: Cytosine methylation
FILE REFERENCE: 5013.1014
CURRENT APPLICATION NUMBER: US/10/311,455
CURRENT APPLICATION NUMBER: PCT/EP01/07537
PRIOR APPLICATION NUMBER: PCT/EP01/07537
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US-10-424-599-59036/c
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US-10-311-455-1419/c
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Publication No. US20040031072A1
GENERAL INFORMATION:
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SEQ ID NO 92893
LENGTH: 284
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SEQ ID NO 59036
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Best Local
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid
TITLE OF INVENTION: Plants and Uses
FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
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LOCATION: (1)..(284)
OTHER INFORMATION: unsure at
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Similarity 95.0%;
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Pred. No. 2.8e
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Pred. No. 2.6e+02;
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Best Local S
Matches 19
                                Sequence 11, Application US/10433793 Publication No. US20040142334A1 GENERAL INFORMATION:
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SEQ ID NO 191
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                                                                                                                                                                                                                               Query Match
Best Local
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2000-03-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 5013.1004
CURRENT APPLICATION NUMBER: US/10/221,613
CURRENT FILING DATE: 2002-09-13
PRIOR APPLICATION NUMBER: PCT/EP01/02945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with Cell Cycle
FILE REFERENCE: 5013.1004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 2424
APPLICANT: Epigenomics AG
TITLE OF INVENTION: Diagnose von mit Anglogenese assoziierten Krankheiten
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: OLEK, ALexander APPLICANT: PIEPENBROCK, C
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DE 10019058.8
DE 10019173.8
DE 10032529.7
DE 10043826.1
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                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: chemically treated genomic DNA
                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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APPLICATION NUMBER: DE 10032529.7
FILING DATE: 2000-06-30
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0; Mismatches
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Pred. No. 3e+C
O; Mismatches
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3e+02;
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FILE REFERENCE:

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FEATURE:
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(54016)
COTHER INFORMATION: n = A,T,C or G, or insertion/deletion polymorphism (see Tables 1-US-10-741-600-17886)
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US-10-741-600-17886/c

US-10-741-600-17886/c

Sequence 17886, Application US/10741600

Publication No. US200S0026169A1

GENERAL INFORMATION:

APPLICANT: CARGILL, Michele et al.

TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF DETECTION AND USES THEREOF FILE REFERENCE: CL001499

CURRENT APPLICATION NUMBER: US/10/741,600

CURRENT FILING DATE: 2003-12-22

NUMBER OF SEQ ID NOS: 73997

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 17886

LENGTH: 54016

TYPE: DNA

OCCANTENT AND COMMENTED OF THE COMMENT OF TH
Search completed: September Job time: 30.2273 secs
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                                                                                                                                                                                                                                                                                                                                                                                      Query Match 83.6%; Score 18.4; DB 21; Length 54016; Best Local Similarity 95.0%; Pred. No. 3.3e+02; Matches 19; Conservative 0; Mismatches 1; Indels 0;
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CURRENT FILING DATE: 2003-06-06
NUMBER OF SEQ ID NOS: 212
SEQ ID NO 11
LENGTH: 23695
TYPE: DNA
ORGANISM: Artificial Sequence
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Perfect score:
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Match Length
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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Gapop 10.0 , Gapext 1.0
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US-09-280-181B-2
US-08-781-966A-1722
US-08-781-966A-1722
US-09-949-016-87851
US-09-949-016-186785
US-09-949-016-186785
US-09-949-016-186786
US-08-956-171E-38
US-08-956-171E-38
US-08-949-016-17554
US-09-949-016-17554
US-09-949-016-17554
US-09-949-016-17559
US-09-949-016-12368
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US-09-949-016-12598
US-09-949-016-12583
US-09-949-016-1368-50
US-09-939-016-158-50
US-09-339-338-50
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Sequence 38, Appl
Sequence 38, Appl
Sequence 17554, A
Sequence 17554, A
Sequence 17978, A
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Sequence 1722, A
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12505, A
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AT TOWNSHIP	US-09-702-705-1792	US-09-671-317-149	US-09-221-017B-741	US-09-949-016-157860	US-09-949-016-157859	US-09-949-016-157858	US-09-949-016-157857	US-09-658-824-1279	US-09-671-325-1279	US-09-614-124B-1279	US-09-736-457-1279	US-09-702-705-1279	US-09-551-621-50	US-09-590-751A-50	US-09-834-759-50	US-09-285-480-50	US-09-604-287A-50	US-09-433-826B-50
	Sequence 1792, Ap	Sequence 149, App	Sequence 741, App	Sequence 157860,	Sequence 157859,	Sequence 157858,	Sequence 157857,	Sequence 1279, Ap	Sequence 1279, Ap	Sequence 1279, Ap	Sequence 1279, Ap	Sequence 1279, Ap	Sequence 50, Appl					

ALIGNMENTS

RESULT 1 US-09-280-181B-2

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| Sequence 2, Application US/09280181B | Patent No. 6280941 | GENERAL INFORMATION: | APPLICANT: Betty P. Tsao (Inventor) | APPLICANT: Rita M. Cantor (Inventor) | APPLICANT: Jerome I. Rotter (Inventor) | APPLICANT: Jerome I. Rotter (Inventor) | TITLE OF INVENTION: Genetic Marker Test for Lupus | FILE REFERENCE: P07 41735 | CURRENT APPLICATION NUMBER: US/09/280,181B | CURRENT FILING DATE: 1999-03-29 | NUMBER OF SEQ ID NOS: 4 | SOFTWARE: FastSEQ for Windows Version 4.0 | SEQ ID NO 2 | LENGTH: 20 | TYPE: DNA | ORGANISM: Homo sapiens
                                                                                                                                                                                                                                         US-08-950-1.
; Sequence 1722, Appiron
; Sequence 1722, Appiron
; Patent No. 6593114
; GENERAL INFORMATION:
APPLICANT: Charles Kunsch
Gil H. Choi
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             CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences,
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
COUNTRY: USA
ZIP: 20850
                                                                                                                                              Patrick S. Dillon
Craig A. Rosen
Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
COMPUTER READABLE FORM:
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Best Local Similarity
GENERAL INFORMATION:
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TELEPAX: (301) 309-8439
INFORMATION FOR EQ ID NO: 1722:
SEQUENCE CHARACTERISTICS:
                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                             SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                      FILING DATE:
ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM: MSDOS vei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 60/009,861
FILING DATE: January 5, 1996
APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Mark J. Hyman
REGISTRATION NUMBER: 46,789
REFERENCE/DOCKET NUMBER: P8248P1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                   STREET: 9410 Key West Avenue CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LICANT: Charles Kunsch
LE OF INVENTION: Staph
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APPLICATION NUMBER: US/08/956,171E
FILING DATE: 20-Oct-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             57 AACGTGTGGTAATGACTGCA 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18;
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                                                                                                                                                                                                                                                                                                                                   20850
                                                                                                                                                                                                                                                                                                                                                                   Maryland
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STRANDEDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 325 base pairs
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                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                                                                                      Human
                                                                                                                                                                                                                                                                                                Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84.0%;
                                                                                                                                                                                                                                                              MSDOS version 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                 Genome Sciences, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Staphylococcus aureus Polynucleotides and Sequences
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                                                                                                                                                                                                               US/08/781,986A
                                                                                                                                                                                                                                                                                              3.50 inch, 1.4Mb storage
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Pred. No. 41;
0; Mismatches
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APPLICANT: VUENTIER, J. Craig et al.
APPLICANT: VUENTION: POLYMORPHISMS IN KNOWN GENES ASS
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING D
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US-09-949-016-87852/c
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US-08-781-986A-1722
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                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR PILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
                                                              NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 87852
LENGTH: 601
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GENERAL INFORMATION:
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ORGANISM: Human
ORGANISM: Human
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TYPE: nucleic acid
STRANDEDNESS: double
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90.0%;
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90.0%; Pred. No. 41;
tive 0; Mismatches
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Pred. No. 46;
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OF DETECTION AND USES THEREOF
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PATENT NO. 6812339;

GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AS
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: 00/241,755
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/247,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FRASTSEQ for Windows Version 4.0
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                                                                                      ; ORGANISM: Human
US-09-949-016-186785
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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Best Local Sim
Matches 18;
Query Match
Best Local Similarity
Matches 18; Conserv
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SEQ ID NO 183202
LENGTH: 601
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 186785,
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                                                                                                                               TYPE: DNA
                                                                                                                                               LENGTH: 601
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Similarity 90.0%;
18; Conservative
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90.0%;
    ; Score 16.8; D; Pred. No. 46; O; Mismatches
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Pred. No. 46;
0; Mismatches
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Pred. No. 40
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                                           Length 601;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 186786
LENGTH: 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 6593114
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 38,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 186786, A Patent No. 6812339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                   FILING DATE: 20-Oct-1997
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                         OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/956,171E
                                                                                                                                                                                  ZIP: 20850
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
                                                                                                                                                                                                                                                                                                                                                                                        Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         392
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APPLICATION NUMBER: 60/009,861 FILING DATE: January 5, 1996 APPLICATION NUMBER: 08/781,986
                                                                                                                                                                                                                                                                            CITY: Rockville
STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAATTGTGGTGATGGCTGCA 373
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                                                                                                                                                                                                                                                             COUNTRY: USA
                                                                                                                                                                                                                                                                                                                   STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/08956171E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Charles Kunsch
Gil H. Choi
                                                                                                                                                                                                                                                                                                                                                                                                                                             Patrick S. Dillon
Craig A. Rosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90.0%;
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                                                                                                                                                                                                                                                                                                                                    Genome Sciences,
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Pred. No. 4
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FILING DATE: January 3, 1997 ATTORNEY/AGENT INFORMATION: NAME: Mark J. Hyman

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US-08-781-986A-38
                                                                                              US-08-781-986A-38
                                                                                                                                                                                        NAME: Benson, Bob
REGISTRATION NUMBER: 30,446
REFERENCE/DOCKET NUMBER: PB24
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEPAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 38:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 38, Application US/08781986A Patent No. 6737248 GENERAL INFORMATION:
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                              Matches
                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: HP Vectra 486,
OPERATING SYSTEM: MSDOS
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (240) 314-12;
TELEPAX: (301) 309-8439
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 23439 base pairs
TYPE: nucleic acid
TYPE: Acid
                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                          FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 9410 Key West Avenue CITY: Rockville
                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                              TOPOLOGY:
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REFERENCE/DOCKET NUMBER: PB248P1
TELECOMMUNICATION INFORMATION:
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                              18;
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AAATTGTGGTAATGACTGCA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 23439 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Maryland
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Charles Kunsch
IVENTION: Staphylococcus aureus Polynucleotides and Sequences
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                                                                                                              linear
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                                            84.0%;
90.0%;
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                            Score 16.8; D
Pred. No. 87;
0; Mismatches
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Pred. No. 87;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     38:
                                                           DB 4;
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                                                            Length 23439;
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RESULT 11
US-09-949-016-17554/c
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                                                                                                                                                            ; ORGANISM: Human
US-09-949-016-11945
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APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT PILICATION NUMBER: 09/29/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FREUENCE for Windows Version 4.0

SEQ ID NO 17554
                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
CURRENT FILING DATE: 2000-04-14
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Matches
                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11945
LENGTH: 86439
                                                                        Matches
                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (1)...(62908)
OTHER INFORMATION: n =
                                                                                                                                                                                                       TYPE: DNA
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ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 62908
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o. 6812339
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AAATTGTGGTAATGACTGCA 20
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                                                                        Conservative
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                                                                      84.0%; Score 16.8; DB 4;
90.0%; Pred. No. 1.1e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 16.8; DB 4; Length 62908; Pred. No. 1e+02; 0; Mismatches 2; Indels 0;
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                                                                                                                 Length 86439;
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23617 AAATTGTGGTGATGATTGCA 23598

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                                                                                                                                                         ; NAME/KEY: misc_feature
; LOCATION: (1)...(123463)
; OTHER INFORMATION: n = A,T,C or
US-09-949-016-17078
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US-09-949-016-16990
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
                                                                          Query Match
Best Local Similarity
Matches 18; Conserv
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17078
LENGTH: 123463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16990, Application US/09949016 Patent No. 6812339
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Patent No. 6812339
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                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                      ORGANISM: Human FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23617 AAATTGTGGTGATGATTGCA 23598
29756 AAATTGTGGTGATGGCTGCA 29737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
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                      AAATTGTGGTAATGACTGCA 20
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                                                                            Conservative
                                                                                         84.0%;
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90.0%;
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                                                                          Score 16.8; D
Pred. No. 1.2e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.8; DB 4;
Pred. No. 1.1e+02;
                                                                                         ..2e+02;
                                                                                                                DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 86440;
                                                                                                                Length 123463;
                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                          Gaps
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Search completed: September Job time: 7.26155 secs

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 12257, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CL001307
                                                                       Matches
                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 12257
                                                                                                                                                         TYPE: DNA
ORGANISM: Human
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(129327)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/237,768
                                                                                                                                                                                                                                                                        LENGTH: 129327
68053 AAATTGTGGTGATGGCTGCA 68034
                                                                     18; Conserv
                    1 AAATTGTGGTAATGACTGCA 20
                                                                       Conservative
                                                                                    84.0%;
                                                                                                                                                                                                                                                                                                                              207012
                                                                     0; Mismatches
                                                                                                                                                               ç
                                                                                    Score 16.8;
Pred. No. 1.
                                                                                                                                                               G
                                                                                        .2e+02
                                                                                                          B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ASSOCIATED
OF DETECTION AND USES THEREOF
                                                                                                     Length 129327;
                                                                       Indels
                                                                       0
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Database
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Maximum DB seq length: 200000000
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OM nucleic - nucleic search, using sw model
Published Applications NA:*

1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

6: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

7: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

10: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

11: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

12: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

13: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

15: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

16: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

16: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

18: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

19: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

20: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

21: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

22: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

23: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

24: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

25: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

26: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IDENTITY_NUC Gapop 10.0 , Gapext 1.0
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	n	0	n	n	Ω		Regult No.
7	o	u	•	w	N	_	
16.8	16.8	16.8	18	18.4	20		Score
84.0	84.0	84.0	90.0	92.0	100.0	100.0	Query
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10	20	20	19	21	11	11	BB
US-09-764-891-9678	US-10-719-993-16988	US-10-719-993-16987	US-10-322-281-560	US-10-774-355A-643	US-09-909-317-5	US-09-909-317-2	ID
			Sequence 560, App	Sequence 643, App		Sequence 2, Appli	escription

ALIGNMENTS

US-09-909-317-2

```
Sequence 2, Application US/0999317

Publication No. US20040152075A1

GENERAL INFORMATION:

APPLICANT: Betty P. Tsao (Inventor)

APPLICANT: Rita M. Cantor (Inventor)

APPLICANT: Rita M. Cantor (Inventor)

APPLICANT: Rita M. Canter (Inventor)

APPLICANT: Parome I. Rotter (Inventor)

ITILE OF INVENTION: Genetic Marker Test for Lupus

FILE REFERENCE: 18810-82152

CURRENT PILING DATE: 2001-07-18

PRIOR APPLICATION NUMBER: US/09/909,317

CURRENT FILING DATE: 1999-03-29

NUMBER OF SEQ ID NOS: 7

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 2

LENGTH: 20

TYPE: DNA

ORGANISM: Homo Sapiens

US-09-909-317-2

Query Match

Best Local Similarity 100.0%; Score 20; DB 11; Length 20;

Best Local Similarity 100.0%; Pred. No. 7.1;

Matches 20; Conservative 0; Mismatches 0; Gaps

Qy 1 AAATTGTGGTAATGACTGCA 20

Qy 1 AAATTGTGGGTAATGACTGCA 20

LENGTH: 100.0%; Pred. No. 7.1;

Matches 20; Length 20; Caps
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US-09-909-317-5/c

RESULT

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APPLICANT: ETRESCEI, Stuart
APPLICANT: Zhang, Xinmin
TITLE OF INVENTION: MOUSE OLFACTORY RECEPTOR GENE SUP
FILE REFERENCE: A34570-PCT-USA-A 070050.2520
CURRENT APPLICATION NUMBER: US/10/774,355A
CURRENT FILING DATE: 2004-02-06
PRIOR APPLICATION NUMBER: PCT/US02/2556
PRIOR APPLICATION NUMBER: 60/311,159
PRIOR APPLICATION NUMBER: 60/311,159
PRIOR APPLICATION NUMBER: 60/311,159
PRIOR FILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-08-09
SOFTWARE: FASTSEQ ID NOS: 2596
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NOS 643
LENGTH: 945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/09909317
Publication No. US20040152075A1

GENERAL INFORMATION:
APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Rita M. Cancor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
TITLE OF INVENTION: Genetic Marker Test for
FILE REFERENCE: 18810-82152
CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 7
SECTIMARE: FastSEQ for Windows Version 4.0
RESULT 4
US-10-322-281-560/c
is Sequence 560, Application US/10322281
publication No. US20040126762A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-74-355A-643/c
US-10-774-355A-643/c
; Sequence 643, Application US/10774355A
~~~~11~ation No. US20050043513A1
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                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: MOR204-18 US-10-774-355A-643
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-909-317-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 643, Application No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 5
LENGTH: 2085
                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                       Local
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                                                                                                                                                                                                                   19;
                                                                                                                                                                          1 AAATTGTGGTAATGACTGCA 20
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                                                                                                                                                                                                                                       Similarity
                                                                                                                                          AAATTGTGGTAATGACTACA 594
                                                                                                                                                                                                                     Conservative
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100.0%; Pred. No.
                                                                                                                                                                                                                                       92.0%;
95.0%;
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                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                       Score 18.4;
Pred. No. 79;
                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 2085;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                         Length 945;
                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                   Gaps
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; FEATURE:
, NAME/KEY: misc feature
; LOCATION: (1)...(68732)
; OTHER INFORMATION: n = A,T,C or
US-10-322-281-560
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Homo sapiens US-10-719-993-16987
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                                                                                                                                                                                                                                                                                                                                                                                          RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 5
                                                                                                                                                                                                                                                                                                                                                                     US-10-719-993-16988/c
US-10-719-993-16988
                                                                                                                                                                                                     Sequence 16988, Application US/10719993

Publication No. US20040265849A1

GENERAL INFORMATION:

APPLICANT: CARGILL, Michele et al.

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH

TITLE OF INVENTION: ALZHEIMER'S DISEASE, METHODS OF DETECTION

FILE REFERENCE: CL001496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/719,993
CURRENT FILING DATE: 2003-11-24
NUMBER OF SEQ ID NOS: 55342
SOFTWARE: FAStSEQ for Windows Version 4.0
SEQ ID NO 16987
LENGTH: 201
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16987, Application US/10719993

Publication No. US20040265849A1

GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.

TITLE OF INVENTION: ACCHEINCE OLLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: ALCHEIMER'S DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001496
                                          CURRENT APPLICATION NUMBER: US/10/719,993
CURRENT FILING DATE: 2003-11-24
NUMBER OF SEQ ID NOS: 55342
SOFTWARE: FAStSEQ for Windows Version 4.0
SEQ ID NO 16988
LENGTH: 201
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 90.0%; Score 18;
Best Local Similarity 100.0%; Pred. No.
Matches 18; Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 560
LENGTH: 68732
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 84.0%; Score 16.8; DB 20
Best Local Similarity 90.0%; Pred. No. 3.7e+02;
Matches 18; Conservative 0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: David W. Morris
APPLICANT: Marc S. Malandro
TITLE OF INVENTION: NOVel Compositions and Methods
FILE REFERENCE: 529452001000
CURRENT FAPPLICATION NUMBER: US/10/322,281
CURRENT FAPPLICATION NUMBER: 002-12-17
CURRENT FILING DATE: 2002-12-17
NUMBER OF SEQ ID NOS: 866
                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13183 ATTGTGGTAATGACTGCA 13166
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2.4e+02;
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RESULT 8
US-08-781-986A-1722/c
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US-09-764-891-9678
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; ORGANISM: Homo sapiens
US-09-764-891-9678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 10231
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 9678
LENGTH: 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1722, Application US/08781986A
Publication NO. US20030054436A1
GEMERAL INFORMATION:
APPLICANT: Charles Kunsch
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Best Local Similarity
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NAME: Benson, Bob
REGISTRATION NUMBER: 30,446
REFERENCE/DOCKET NUMBER: PB248PP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEPHAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 1722:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/764,891
CURRENT FILING DATE: 2001-01-17
                                                                                                                                                                                                                                 OPERATING SYSTEM: MSDO
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences NUMBER OF SEQUENCES: 5255
                                                                                                                                                                  APPLICATION NUMBER: 1
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 5:
CORRESPONDENCE ADDRESS:
                                                                                                                ATTORNEY/AGENT INFORMATION:
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                                                                                                                                   APPLICATION NUMBER: FILING DATE:
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9410 Key West Avenue
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90.0%;
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Pred. No. 3.8e+02;
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SEQUENCE DESCRIPTION: SEQ ID NO: US-10-329-624-1722
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                                                     Matches
                                                               Query Match
Best Local Similarity
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LENGTH: 325 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                 SOFTWARE: ASCII Text
CURRENT APPLICATION NUMBER: US/10/329,624
PILLING DATE: 27-Dec-2002
PRIOR APPLICATION NUMBER: 08/956,171
APPLICATION NUMBER: 08/956,171
FILLING DATE: October 20, 1997
APPLICATION NUMBER: 60/009,861
FILLING DATE: January 5, 1996
APPLICATION NUMBER: 08/956,171
FILLING DATE: January 3, 1997
AFTORNEY, AGENT INFORMATION:
NAME: MAIK J. Hyman
REGISTRATION NUMBER: 46,789
REFERENCE/DOCKET NUMBER: PB248P1D1
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MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides
NUMBER OF SEQUENCES: 5256
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ADDRESSEE: Human Genome Sciences, Inc
                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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1 AAATTGTGGTAATGACTGCA 20
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18; Conserv
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                                                                                                                                                                   LENGTH: 325 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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ZIP: 20850
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                      TELEPHONE: (240)
TELEFAX: (301) 30
                                                     Conservative
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Craig A. Rosen
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                                                                   90.0%;
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                                                                    Score 16.8;
Pred. No. 4
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                                                     Mismatches
                                                                                                                                                                                                                                                                                                         PB248P1D1
                                                                                                                                      1722:
                                                                    4e+02;
                                                                                      DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8; Length 325;
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                                                                                  Length 325;
                                                     Indels
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                                                     Gaps
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AACGTGTGGTAATGACTGCA 38

US-10-027-632-218409/c

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APPLICANT: Wang, David G.
APPLICANT: Wang, David G.
TITIE OF INVENTION: Identification and Mapping of Single Nucleotide
TITIE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR PILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Wang, David G.
FITTLE OF INVENTION: Identification and Mapping of Single Nucleotide
FITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108027.129
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/193,483
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US-10-027-632-218409
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____ rocal Similarity
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SEQ ID NO 218409
LENGTH: 663
SEQ ID
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SOFTWARE:
                                                                                PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
                                                           NUMBER OF SEQ ID NOS: 325720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
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                             FastSEQ for Windows Version 4.0
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90.0%;
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Pred. No. 4.5e
0; Mismatches
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_18914C.1 US-10-424-599-53048
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Clone ID: 700605169_FLI US-10-425-114-6765
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                                                                                                                                       TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 53048
LENGTH: 2591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 6765
LENGTH: 1386
                                                                                                                                                                                                                                                                                                       Sequence 53048, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local Similarity
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 Query Match
Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                      APPLICANT: Kovalic Dav. APPLICANT: Zhou Yihua APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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ORGANISM: Glycine max
FEATURE:
                                                                                                         TYPE: DNA
ORGANISM: Glycine
                                                                                         FEATURE:
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Kovalic, David K.
Screen, Steven E
Tabaska, Jack E
Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAATTGTGGTAATGACTGCA 20
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o. US20040034888A1
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90.0%;
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Pred. No. 4.5e+02
0; Mismatches
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Score 16.8; DB 18
Pred. No. 5.5e+02;
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                  DB 18;
                      Length 2591;
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RESULT 15
US-10-329-624-38
; Sequence 38, Application US/10329624
; Publication No. US20040043037A1
; GENERAL INFORMATION:
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US-08-781-986A-38
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Publication No. US20030054436A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY AGENT INFORMATION:
NAME: Benson, Bob
REGISTRATION NUMBER: 30,446
REFERENCE/DOCKET NUMBER: PB24
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE:
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MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33

OPERATING SYSTEM: MSDOS version 6.2

OPERATING SYSTEM: MSDOS version 6.2
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TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 9110
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 23439 base pairs
TYPE: nucleic acid
STRANDEDESS: double
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                                                                                                                                                                                                                                                                                                                  22929 AACGTGTGGTAATGACTGCA 22948
                                  Patrick S. Dillon
Craig A. Rosen
Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus aureus Polynucleotides and Sequences
NUMBER OF SEQUENCES: 5256
                     CORRESPONDENCE
                                                                                                                                                               APPLICANT: Charles Kunsch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2288 AAATTGTGGTTATGATTGCA 2307
                                                                                                                                                                                                                                                                                                                                                                                           18; Conservative
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Y: USA
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ADDRESSEE: Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Human Genome Sciences, Inc.
9410 Key West Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                Gil H. Choi
                       ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                       84.0%; Score 16.8; DB 8; Length 23439;
90.0%; Pred. No. 7.8e+02;
Live 0; Mismatches 2; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US/08/781,986A
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Search completed: September Job time: 26.1157 secs
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                                                                                                                                                            Matches
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Best Local &
                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 38: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/956,171
FILING DATE: October 20, 1997
APPLICATION NUMBER: 60/009,861
FILING DATE: January 5, 1996
APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Mark J. Hyman
REGISTRATION NUMBER: 46,789
REGISTRATION NUMBER: 46,789
                                                                               22929 AACGTGTGGTAATGACTGCA 22948
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
COMPUTER: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
COMPATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/329,624
                                                                                                                                                                                                                                                                                                                                                                                                  TELEPAX: (301) 309-8439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 27-Dec-2002 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                          l Similarity
18; Conserv
                                                                                                                     1 AAATTGTGGTAATGACTGCA 20
                                                                                                                                                                                                                                                                               LENGTH: 23439 base pairs TYPE: nucleic acid STRANDEDNESS: double TOPOLOGY: linear
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CITY: Rockville
STATE: Maryland
COUNTRY: USA
                                                                                                                                                          Conservative
                                                                                                                                                          84.0%; Score 16.8; DB 18; 90.0%; Pred. No. 7.8e+02; tive 0; Mismatches 2;
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Result
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Maximum
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
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; TYPE: DNA
; ORGANISM: Human
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; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(98439)
; OTHER INFORMATION: n = A,T,C or
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PALENT NO. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN G.
TITLE OF INVENTION: WITH HUMAN DISEASE, MET FILE REFERENCE: CLO01307
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-3
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR PILING DATE: 2000-09-08
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SOFTWARE: FRStSEQ for Windows Version
SEQ ID NO 13597
LENGTH: 98439
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                                                         TTTCCACCATCCAGAAGCAACTGGTGTTAATATCTGGAACATAAAATGGAATCACATCTT 37049
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WS-09-949-016-12360
US-09-949-016-12360
; Sequence 12360, Application US/09949016
; Sequence 12360, Application US/09949016
; Patent No. 6812339
; Patent No. 6812339
; Patent INFORMATION:
APPLICATION: FOLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
TITLE REFERENCE: CL001307
; FILE REFERENCE: CL001307
; CUGRENT APPLICATION NUMBER: US/09/949,016
; CUGRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR APPLICATION NUMBER: 60/241,756
; PRIOR APPLICATION NUMBER: 60/237,768

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RESULT 4
US-09-949-016-16409
; Sequence 16409, Application US/09949016
; Patent No. 6812339
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US-09-949-016-12360
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PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 12360
LENGTH: 105679
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 10.9%;
Best Local Similarity 60.6%;
Matches 452; Conservative
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 16409
LENGTH: 107679
TYPE: DNA
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Best Local Similarity
Matches 452; Conserv
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
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AGCCTGGGCAACATGGAGAGATTCCC 831
                                                           AATCCCAGCAATTTGGGAAGTCGAGGTGGGAGGATTGCTTGAGGCCAGGGGTTCAAGACC
                                                                                              TAAATGCAACATCCTTTAGATAAAAATATTTGTCGGCCGGGCGCGGTGGCTCACGCCTGT
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Pred. No. 7.2e-44;
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GENERAL INFORMATION:
APPLICANT: PXE International, Inc.
APPLICANT: University of Hawaii
TITLE OF INVENTION: Mutations in a gene enco
TITLE OF INVENTION: Pseudoxanthoma Elasticu
FILE REFERENCE: PXE-001
CURRENT APPLICATION NUMBER: US/09/792,616
CURRENT FILING DATE: 2001-02-23
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.0
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US-09-792-616-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 1
LENGTH: 107820
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE: misc feature
NAME/KEY: misc feature
OTHER INFORMATION: "n" can be an A or a T or a G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 440; Conserv
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AAACAACAAGATCTAGGCCAGGTGCAGTGGCTCATGCCTGTAATCCCAGCAATTTGGGAA 764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGTGATCTCAGCTCACTGCAACCTCCAGCTTCAAGTGATTCTCCTGCCTCAGC
                                                 CATATGCATGAGCATTCGAGTTTTTTTTGGAATATTCAGCAATTACAATCAAAAATGCCAT
                                                                                 CATAGGATTAGAAAAGAAACCAATTTTAATAAACTATATTTTGAAGTATAGTTCTATATT
                                                                                                                                                ACACTTTTTATGAGTTTTATTCATCTGCTTATTTTTCTTATCCTCTATACCAGAATTGAA 589
                                                                                                                                                                                                                                             CACGTTCTTCCTCTCTCTCTTTCTTCCTAATGATGACATTTATGTACTGCACATGA
                                                                                                                                                                                                                                                                    AACCTTCTTGCTTACTCAACATCCTTGTCTCTTAAGCCACTAGCTTCTTCTCTATGGTTA
                                                                                                                                                                                                                                                                                                          TGGTCTCATCTGATTTTTAGCAACGGTGGGGTCAACCTGATCAATTGCCCCTGAATTTTCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTCTCTAGTAGCTGGGATTACAGGCACATGCCACCATGCCCCGGCTAATTTTTATATTTTC 7483
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                                                                                                                CTGGAATACAAGGTGAAGACACTGGAAAAAGTCCTGGCCCCTAGCCCTAGCAAAAACAAT
                                                                                                                                                                              AGCCCTGTCTGAGTGCCTTATGTTGTAATTGCTAAGCACCTGCCTATTGTTTGCAAGGCC
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; ORGANISM: Human
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)...(36311)
; OTHER INFORMATION: n = A
US-09-949-016-13627
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US-09-949-016-13627/c
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SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 13627
LENGTH: 36311
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Best Local Similarity
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-03

PRIOR PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: 60/231,498

PRIOR PRI
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ACATCCTTGTCTCTAAGCCACTAGCTTCTTCTCTATGGTTAACA-----
                                                                        ATAATCACTCCAACCTCTTCCCCTTTATTCCCCCAGCCCCAGGAATTGTAGCTGCTTCCTCC 9141
                                                                                                                                                                                                                      AAGTGCTAGGATTACAAGCGTGAGCCACTGCGCCCCGGCCATTCTTCCAAGTTTCCTGGCT
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                                                                                                                                             GTTACTAACTTTTAGGATTCGGCCTATGTCTCACAACC-----TTCTTGCTTACTCA 487
                                                                                                                                                                                                                                                                                           AAGTGCTAGGATTATAGGCATGAGCCACTGTGCCCAGCCTACCTTCAACGTATCTAACTG
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Pred. No. 4e-43;
0; Mismatches 5
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  GGCGGAGCTTGCAGTGAGCCGAGATTTCGCCACTGCACTCCAGCCTGG
                                                                                                                               GAGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGTGGGAGGATCGC-TGGGCTCAGGA
                                                                                                                                                                                             AACTGTGGGAGGCTGAGGCAGGTGGATCACAAG-GTCAGGAGATCGAGACCATCCTGGCT
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                                              GGACGCCTGTGGTCCCAGCTACTCAGGAGGCTGAAGTGGGAGAATGGCATGAACCCGGGA
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 15714
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISSASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
FILE REFERENCE: CL001307
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Best Local (
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TYPE: DNA
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ACAACAAGATCTAGGCCAGGTGCAGTGGCTCATGCCTGTAATCCCAGCAATTTGGGAAGT 766
                                    AÇAACÇITCTTGCTTA-ÇTCAAÇATCCTTGTÇTCTTAAGÇÇACTAGCTTCTTCTCTATGG 526
                                                                                                                                                                                                                                                                                                                     CCAG-CCTACCTTCAACGTATCTAACTGGTTACTAACTTTTTAGGATTCGGCCTATGTCTC
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Pred. No. 4.6e-43;
0; Mismatches 357;
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. CRAIG et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-09-08
NUMBER: 00 520 ID NOS: 207012
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 12032
LENGTH: 40655
TYPE: DNA
ORGANISM: Human
US-09-949-016-12032
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Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J.
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Best Local Similarity 52.8%;
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                               AACATCCTTGTCTCTTAAGCCACTAGCTTCTTCTCTATGGTTAACACTTTTTATGAGTTT 546
                                                                       TTTTTATTATCAAAACAATATTATGTTACGAAAA----
                                                                                                      ATCTAACTGGTTACTAACTTTTAGGATTCGGCCTATGTCTCACAACCTTCTTGCTTACTC
                                                                                                                                          GGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCGTGCCCGGCTTA----GAACAT
                                                                                                                                                                  AGCCTCCCAAAGTGCTAGGATTATAGGGCATGAGCCACTGTGCCCCAGCCTTCAACGT
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                                                                                                                                                                                                                                                                                    CCTGCCACCACCCCAGCTAATTTTT-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                GTCTTACTCTGTCACCCAGGCTGGAGTGCAGTGGCTCAATCTCGGGTCACTGCAGCCTCC
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-0-08
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
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US-09-949-016-15919
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GENERAL INFORMATION:
APPLICANT: VENTER, J.
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Best Local Similarity 52.8%;
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                                                                                                                                                       GAGGATTGCTTGAGCCCAGGAGATTGAGACTGCAGTAAGCCAAGGTTGCACCATTGCACT
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RESULT 10
US-09-949-016-14577
; Sequence 14577, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTICR, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755

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; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSEQ for Windows Version 4.
; SEQ ID NO 14577
LENGTH: 678533
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, NAME/KEY: misc_feature
; LOCATION: (1)...(678533)
; OTHER INFORMATION: n = A,T,C
US-09-949-016-14577
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Matches 716
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ATCCGGTGAAACCTCGTCTCTACTAAAAATACAAAAAATTAGCCGAGCGTGGTGGCGGG 405916
                                   CTTTGGGAGGCCGAGGCGGGCGGAT---CACGAGGTCAGGAGATCGAGACCATCCTGGCTA
                                                                                               ATTTGGGAAGTCGAGGGTGGGAGGATTGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCA
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Pred. No. 1.5e-41;
0; Mismatches 662;
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RESULT 11
US-09-949-016-14578
; Sequence 14578, Application US/09949016
; Patent No. 6812339
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                                TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF FILE REFERENCE: CLO01307

CURRENT FILING LOTE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-20

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-01-03

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: PASTSEQ for Windows Version 4.0

SEQ ID NO 14578

LENGTH: 678533
                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
ORGANISM: Human
                      TYPE: DNA
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; FEATURE: ; NAME/KEY: misc feature ; LOCATION: (1). - (678533) ; OTHER INFORMATION: n = A,T,C US-09-949-016-14578

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Best Local Similarity
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TTTGCTCATGCTATTTTCATTGCCTGGAGTGTCTTCTCTCTGGATAGAAAAATCCTTTTCC
                                  CTAGGTTTCAGTTGGTATTTTGTCCCGACGG----TCTGTGGACGGCAGGTTAGAACGC 1110
                                                                                                      AAAGTCACAGGTACTGCTAATACTCCTGGTATTTGTAGTAAATTCATAATAAAGGAAATG
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Pred. No. 1.5e-41;
O; Mismatches 662;
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; ORGANISM: Human
US-09-949-016-12513
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US-09-949-016-12513/c
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CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 12513
LENGTH: 103987
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12513, Application US/09949016 Patent No. 6812339
                                                                                                                                                                                                                                                                                     Matches 426;
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                                                                  TGAGTGGCTGGGACTACAAGCACACGCCACGCCTGGTTAATTTGTTGTATTTTTAGT
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                  AGAGACAAGGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAGGGATCC
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                                                                                                                                                                                                                                                                                 Score 219.2; DB 4;
Pred. No. 9.1e-42;
0; Mismatches 283;
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT PILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-39
PRIOR FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FRASESEQ for Windows Version 4.0
LENGTH: 103988
TYPE: NYA
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US-09-949-016-17050/c
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Matches
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GENERAL INFORMATION:
                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                           Local Similarity
                                                               48509
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Conservative
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                                                                                                                                       Score 219.2; DB 4;
Pred. No. 9.1e-42;
0; Mismatches 283;
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                                                                                                                                                                           Length 103988;
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR PILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
VOMBER OF SEQ ID NOS: 207012
SOPTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 12086
LENGTH: 152582
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US-09-949-016-12086
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                                                                                                                                                                                                                                                                                                                                               Sequence 12086, Application US/09949016
PACENT NO. 6812339
GENERAL INFORMATION: APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION FILE REFERENCE: CL001307
NAME/KEY: misc_feature
LOCATION: (1)...(152582)
                                                              TYPE: DNA
ORGANISM: Human
                                              FEATURE:
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Sequence 17390, Application US/09949016

PATENT NO. 6812339

GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASS

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF

FILE REFERENCE: CL001307

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                      RESULT 15
US-09-949-016-17390
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Best Local Similarity
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PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION UNMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.
SEQ ID NO 17390
LENGTH: 152583
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; NAME/KEY: misc feature
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Copyright (c) 1993 - 2005 Compugen Ltd.
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/ Cgn2_6/ptodata/2/pubpna/US07_NEW PUB.seq:*
/ Cgn2_6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
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/ Cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
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                                                                                                                                                        . 6/ptodata/2/pubpna/US11Ā PUĒCOMB.seq:
. 6/ptodata/2/pubpna/US11Ā PUB.seq:*
. 6/ptodata/2/pubpna/US60Ā PUB.seq:*
. 6/ptodata/2/pubpna/US60ĀPUB-COMB.seq:*
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SUMMARIES
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06 NEW PUB.seq:*
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100.0 58.7 58.7 58.7 58.7 57.7	Query Match
2085 10619 10619 10619 10619 10619 10619	
11 15 15 16 17	BG
11 US-09-909-317-5 14 US-10-239-676-1 15 US-10-211-455-43 15 US-10-240-453-1 18 US-10-240-589C-1 14 US-10-239-676-2 15 US-10-311-455-44	ID
Sequence 5, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 2, Appli Sequence 2, Appli Sequence 44, Appl	Description

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	10.9 10.9 10.6 10.5	11.7 11.7 11.7 11.5 11.3		57.7 57.7 112.5 11
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US-10-723-860-2768 US-10-981-277-35 US-10-322-696-70 US-10-741-600-17761	792-616-1 764-328-1 764-328-1 484-577-661 997-722-238 997-723-860-232 723-860-232 074-024-446	US-10-212-872-986 US-10-212-872-987 US-10-084-817-316 US-09-960-253-106 US-09-960-253-100 US-10-334-143-100 US-10-741-600-17781	-10-363-345A -10-363-483A -10-363-483A -10-363-483A 09-764-860-98 09-764-860-98 09-764-860-98	US-10-240-453-2 US-10-240-589C-2 US-10-027-632-154183 US-10-027-632-154183 US-09-960-253-107 US-09-960-253-107 US-09-844-87-300 US-10-103-587A-3 US-10-163-587A-3 US-10-163-587A-3 US-10-163-345A-33309 US-10-363-345A-33309 US-10-363-345A-33300 US-10-363-345A-33300 US-10-363-345A-11999
Sequence 2768, Ap Sequence 35, Appl Sequence 70, Appl Sequence 17761, A	σ	Sequence 986, App Sequence 987, App Sequence 316, App Sequence 106, App Sequence 100, App Sequence 17781, A	12000 11999 12000 12000 986, 1 987, 1 987, 1	Sequence 2, Appli Sequence 15, Appli Sequence 154183, Sequence 107, App Sequence 300, Appli Sequence 3, Appli Sequence 3, Appli Sequence 3, Appli Sequence 3329, Ap Sequence 33390, A Sequence 33399, A Sequence 33399, A Sequence 33399, A

ALIGNMENTS

US-09-909-317-5

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                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 2085
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5, Application US/09909317 Publication No. US20040152075A1 GENERAL INFORMATION:
                                                                                Query Match
Best Local Similarity
Matches 2085; Conserv
                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
ITILE OF INVENTION: Genetic Marker Test for Lupus
FILE REFERENCE: 18810-82152
CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
                      1 TTTAGGGATGATATAGTTGTCAACCCAGAGATGGCATGATCATGCCTTTTGACTTGGTCA 60
  TTTAGGGATGATATAGTTGTCAACCCAGAGATGGCATGATCATGCCTTTTGACTTGGTCA
                                                                                  Conservative
                                                                                100.0%; Score 2085;
100.0%; Pred. No. 0;
tive 0; Mismatches
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                                                                                0,
                                                                                                                          Length 2085;
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	1141 TGCAGGGTCCACCTCGGGCCAATCAACTATATTCCCGAGGCGGGGGGCCTGCGCTTCCCGG 1200	
	1081 CGACGGTCTGTGGACGGCAGGTTAGAACGCCCGTCCAAGCCAGGAGGGTGGACCTAGCAC 1140	
.==	1021 CTGGTATTTGTAGTAAATTCATAATAAAGGAAATGCTAGGTTTCAGTTGGTATTTTTGTCC 1080	
	961 CTAACATGAAAACGTCTGTGATGACTATTGCCCACAAAGTCACAGGTACTGCTAATACTC 1020	
· · · · · · · · · · · · · · · · · · ·	901 CCACAATTTCGAGTAGTGATGAGCTTAATAATATTTCGAGTTATCACCAACAACTGTAAA 960	
	841 CTTTACACACACACACACACACACACAAAATATCTGATAGCAACAGGTGCAGTCATTA 900 	
	781 TGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACATGGAGAGATTCCCCATCTCTTT 840	
	721 GCCAGGTGCAGTGGCTCATGCCTGTAATCCCCAGCAATTTGGGAAGTCGAGGTGGGAGGAT 780	
	661 AAACCAATTTTAATAAACTATATTTTGAAGTATAGTTCTATATTAAACAACAAGATCTAG 720	
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	361 CACCTCAGCCTCCCAAAGTGCTAGGATTATAGGCATGAGCCACCTGTGCCCAGCCTACCTT 420	
	301 GACAAGGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAGGGATCCGGC 360	
	241 AGCCTCCCGAGTAGATGGGACTACAGGCACCTGCCACCACCCCCAGCTAATTTTTGTAGA 300	
	181 TGGCACTATCATGGCTCACCACAGCCTCAACCTTCAGGGCTCAGGTGATCCTCCCACTTC 240	
	121 AAATATCTTTTTTTTTTTTTTTGAGACAGGGTCACACTGTCACCCAGGGTAGAGTCCAG 180	
	61 TTCTCTAAGTAAAACTTTTATTTGTTCCATCATATTTTCCACTTATTCTGTTTACCTTCA 120	

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2041 CCGGATGGC	2041	1981 CGCCAAGAGCGGGCCTCTTGCAAGAAATGCAGCGAGGGATCCCCAAGGACTC	1921 CTTTGGCGGCAGCTAGGGGAGGATGGCGGAGTCTTCCGGATAAGCTCTATCGAGTCGAC	1921 CTTTGGCGGCAGCTAGGGGAGGATGGCGGAGTCTTCGGATAAGCTCTATCGAGTCGAC	1861 CACG	Qy 1861 CACGCGAGGCGAGGCGGCAAGCGTGTGTTTTCTAGGTCGTGGCGTCGGGCTTCCGGAG	Db 1801 GTGGCCGGTGCGCGTGTTCGGTGCGCTCTAGCCGTGGCCGGCTGAGCCG	1801	1741	QY 1741 TGGACGCGGGTTCCCGTGGCGTTCCCGCGGGCAGGCATCAGCAATCTATCAGGGAACGGCG		1681	1621 CTAGCTCAGCCCAAGCCCCGCCTCGGCCCCAGGGCAGCGGCCGCAGAGCTCCAC	1621 CTAGCT	Db 1561 AAGTAAGACTTCCTGGGACAGAACAATCAAAGGGGTGGCGCCGGGTCCTCCAAAGAGCT	1561	Db 1501 TGAGACTCAGTCTCAAAAATAAAAGGGGGAGGGGTTGGGGGGTAAAATTAGTTGTGAAAA	1501	Db 1441 CTGGGCTCAGGAGTTCCAGACTGCAGTGAGCCATGATGGCGGCACTGCACTCCAGCGCGC	1441		1381	1321 TTGAGACCAGTCTAGCCAACTTGGCGAGAC	1321	Db 1261 CATGCCCCTGATCCCAGCACTTCGGGAGGCTGAGGCGTGAAGATCACTTGTAGCAGGAG	, 1261 CATGCCC	1201 ACCCAGCTG	Qy 1201 ACCCAGCTGCCCTCAGGGGAGAGAGACACACTTAAGAGTTTGGGGCCGGCGTGGTAGCT	Db 1141 TGCAGGGTCCACCTCGGGCCAATCAACTATATTCCCGAGGCGGGGGCCTGCGGTTCCCG
		GCT 2040	TA 198	ГА 198 	AG 1920	;AG 1920 -	CG 1860	1 CG 1860	CG 1800	CG 1800	CG 1740	CG 1740	GC 1680	GC 1680	TA 1620	TA 1620	TC 1560	TC 1560	GG 1500	.GG 1500	CG 1440	CG 1440	TT 1380	TT 1380	GT 1320	GT 1320	CT 1260	CT 1260	GG 1200

RESULT 2
US-10-239-676-1
US-10-239-676-1
; Sequence 1, Application US/10239676
; Publication No. US20030082609A1
; GENERAL INFORMATION:
; APPLICANT: PIEDENBROCK, Christian
; APPLICANT: BERLIN, Kurt
; APPLICANT: BERLIN, Kurt
; APPLICANT: Diagnosis of Diseases Associated with Gene Regulation

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FILE REFERENCE: 5013.1003
CCURRENT APPLICATION NUMBER: US/10/239,676
CURRENT FILING DATE: 2002-09-24
PRIOR APPLICATION NUMBER: PCT/EP01/03968
DE 10019058 8
DE 10019173.8
DE 10032529.7
DE 10043826.1
PRIOR FILING DATE: 2001-04-06
2000-04-06
2000-04-07
2000-06-30
1000-09-01
NUMBER OF SEQ ID NOS: 228
SEQ ID NO 1
LENGTH: 10619
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US-10-239-676-1
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Best Local Similarity 77.2%;
Matches 1618; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial
FEATURE:
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                                    TTACTCAACATCCTTGTCTCTTAAGCCACTAGCTTCTCTCTATGGTTAACACTTTTTAT
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AAATA--TTTTTTTTTTTTTTTTGAGATAGGGTTATATTGTTATTTAGGTTAGAGTTTAG
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Pred. No. 4.6e-289;
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•	용 성	문 성	dg VQ	94 94	g &	Query Ma Best Loc Matches	; OR ; PE ; OT	SEQ ;	PRIOR		APE	Publ	RESULT	8 &	B 8	8 8	g dy	å S
	241 AGCCTCCCGAGTAGATGGGACTACAGGCACCTGCCACCCCAGCTAATTTTTTGTAGA 300	181 TGGCACTATCATGGCTCACCACAGCCTCAACCTTCAGGGCTCAGGTGATCCTCCCACTTC 240	121 AAATATCTTTTTTTTTTTTTTGAGACAGGGTCACACTGTCACCCAGGCTAGAGTCCAG 180	61 TTCTCTAAGTAAAACTTTATTTGTTCCATCATATTTTCCACTTATTCTGTTTACCTTCA 120	1 TTTAGGGATGATATAGTTGTCAACCCAGAGATGGCATGATCATGCCTTTTGACTTGGTCA 60	y Match Local Similarity 77.2%; Score 1223.6; DB 15; Length 10619; hes 1618; Conservative 0; Mismatches 459; Indels 19; Gaps 10;	ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens) 0-311-455-43	NUMBER OF SEQ ID NOS: 2424 EQ ID NO 43 EQ ID NO 619 TYPE: NA619	OR APPLICATION NUMBER: DE 10032529.7 OR FILING DATE: 2000-06-30 OR APPLICATION NUMBER: DE 10043826.1 OR APPLICATION NUMBER: DE 10043826.1		APPLICANT: BERLIN, Kurt TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Determ TITLE OF INVENTION: cytosine methylation FILE BEFERRACE: 5013 1014	Publication No. US20030143606A1 GENERAL INFORMATION: APPLICANT: OLEARNDEROCK. Christian APPLICANT: PIEPENBROCK. Christian	3 311-455-43 Port 43 April Cation US/10311455	2030 AAGGACTCGCTCCGGATGGCCATCATGGTGCAGGTGCGGGCCGCTGTGCGGGCGG	1970 CGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCTTGCAAGAATGCAGCGAGAGCATCCCC 2029	1910 GGCTTCCGGAGCTTTGGCGGCAGCTAGGGGAGGATGGCGGAGTCTTCGGATAAGCTCTAT 1969	1850 CTGGGTGAGCGCACGCGAGGCGGCGAGGCGGCAAGCGTGTGTTTCTAGGTCGTGGCGTCG 1909	1792 GAAACGGCGGTGGCCGGCGTGTTCGGTGCGCTCTGGCCGTCAGCCGTGGCGG 1849
δ	Db Qy	D S	Q B 5	P B 5	S B &	D 29	Db Qy	Qy Db	φ φ	g Qy	g Qy	Db Qy	Qy Db	Qу	Db Db	Db Qy	D Qy	Db Qy
1374 TITAATTAGCCAGTTGTGGTGAGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGTGGG	1314 CAGGAGTTTGAGACCAGTCTAGCCAACTTGGCGAGACCCTGTCCCTAAAAAAATTTTTTT	72	12	53	11	15 33	955 TGTAAACTAACATGAAAACGTCTGTGATGACTATTGCCCACAAAGTCACAGGTACTGCTA	895 TCATTACCACAATTTCGAGTAGTGATGAGCTTAATAATATTTTCGAGTTATCACCCAACAAC	841 CTTTACACACACACACACACACACAAAATATCTGATAGCAACAGGTGCAG	781 TGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACATGGAGAGAGTTCCCCCATCTCTTT	721 GCCAGGIGCAGIGGCTCAIGCCIGIAATCCCAGCAAITTGGGAAGTCGAGGIGGGAGGAI	661 AAACCAATITTAATAAACTATATTTGAAGTATAGTTCTATATTAAACAACAAGATCTAG 	601 AAAGCACACTCATGTTACAATCTTTGAAATGGAAAAAAAA	541 GAGTTITATTCATCTGCTIATTITICTTATCCTCTATACCAGAATTGAATATTITCAAAT	481 TTACTCAACATCCTTGTCTCTTAAGCCACTAGCTTCTCTCTATGGTTAACACTTTTTAT	421 CAACGTATCTAACTGGTTACTAACTTTTAGGATTCGGCCTATGTCTCACAACCTTCTTGC	361 CACCTCAGCCTCCCAAAGIGCTAGGATTATAGGCATGAGCCACTGTGCCCAGCCTACCTT	301 GACAAGGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAGGGATCCGGC

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Sequence 1, Application US/10240453

Publication No. US20030148326A1

GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: OLEK, Alexander
TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA
TITLE OF INVENTION: Dy Means of Assessing the Methylation Statistic OF INVENTION: With DNA TRanscription

FILE OF INVENTION: With DNA TRanscription

FILE REFERENCE: 5013.1009

CURRENT APPLICATION NUMBER: US/10/240,453

CURRENT APPLICATION NUMBER: DE 10019058.8

PRIOR APPLICATION NUMBER: DE 10019058.8

PRIOR APPLICATION NUMBER: DE 10019173.8

PRIOR APPLICATION NUMBER: DE 10019173.8

PRIOR APPLICATION NUMBER: DE 10032529.7

PRIOR APPLICATION NUMBER: DE 10032529.7

PRIOR APPLICATION NUMBER: DE 10032529.7

PRIOR APPLICATION NUMBER: DE 10043826.1

PRIOR FILING DATE: 2000-04-06

PRIOR APPLICATION NUMBER: DE 10043826.1

PRIOR FILING DATE: 2000-09-01
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SEQ ID NO 1
LENGTH: 10619
TYPE: DNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION: chemically treated
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CGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCTTGCAAGAAATGCAGCGAGAGCATCCCC
                                                                   GGCTTCCGGAGCTTTGGCGGCAGCTAGGGGGAGGATGGCGGAGTCTTCGGATAAGCTCTAT
                                                                                                                          TTGGGTGAGCGTACGCGAGGCGGCGAGGCGGTA---GCGTGTTTTTAGGTCGTGGCGTCG
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APPLICANT: BERLIN, KURT
TITLE OF INVENTION: Diagnosis of Diseases Associated wit
TITLE OF INVENTION: DNA repair
FILE REFERENCE: 5013.1008
CURRENT APPLICATION NUMBER: US/10/240,589C
CURRENT FILING DATE: 2003-09-02
PRIOR APPLICATION NUMBER: PCT/EP01/03972
PRIOR FILING DATE: 2001-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-06
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10043826.1
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; OTHER INFORMATION:
US-10-240-589C-1
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SEQ ID NO 1
LENGTH: 10619
TYPE: DNA
ORGANISM: Artificial Sequence
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APPLICANT: OLEK, Alexander APPLICANT: PIEPENBROCK, Christian APPLICANT: BERLIN, Kurt
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Best Local Similarity
Matches 1618; Conserv
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77.2%;
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Pred. No. 4.6e-289;
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Query Match Best Local Similarity Matches 1600; Conserv

57.7%; ilarity 76.8%; Conservative

Score 1203.2; DB 14; Pred. No. 4.6e-284; 0; Mismatches 463; I

Length 10619;

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APPLICANT: DIEPENBROCK, Christian
APPLICANT: PIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with Gene Regulation
FILE REFERENCE: 5013.1003
CURRENT APPLICATION NUMBER: US/10/239,676
CURRENT FILING DATE: 2002-09-24
PRIOR APPLICATION NUMBER: PCT/EP01/03968
DE 10019058.8
DE 10019058.8
DE 10035829.7
DE 10043826.1
PRIOR FILING DATE: 2001-04-06
2000-04-06
2000-04-07
2000-06-30
2000-09-01
NUMBER: PCT/EP01/03968
; FEATURE:
; OTHER INFORMATION:
US-10-239-676-2
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US-10-239-676-2/c
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                                                           NUMBER OF SEQ ID N
SEQ ID NO 2
LENGTH: 10619
TYPE: DNA
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Publication No. US20030082609A1
GENERAL INFORMATION:
                                           ORGANISM: Artificial Sequence
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              sapiens)
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 TRETTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACATGGAGAGATTCCCCATCTCTTT 840 727 TACTTAAAACCAAAAATTCAAAACCAACACACACAAAAATATCTGATAGCAACAGTCTCTTT 666 841 CTTTACACACACACACACACACACACACAAAAATATCTGATAGCAACAGTCTCTTT 666 842 CTTTACACACACACACACACACACACACACACAAAATATCTGATAACAAACA	
 421 CAACGTARCTAACTGGTTACTAACTTTTAGAATTGGCCTATGTCTCACAACCTTCTGC 480	
 1 TTTAGGGATGATATAGTTGTCAACCCAGAGATGATGATCATGCCTTTTGACCTTTGATCA	

2030 AAGGACTCGCTCCGGATGGCCATCATGGTGGAGGTGCGGGCC 2071	4a VQ
1970 CGAGTCGAGTACGCCAAGAGCGGGGCGCCTCTTGCAAGAAATGCAGCGAGAGCATCCCC 2029	p
1910 GGCTTCCGGAGCTTTGGCGGCAGCTAGGGGAGGATGGCGGAGTCTTCGGATAAGCTCTAT 1969	95 83
1850 CTGGGTGAGCGCACGCGAGGCGGCGAGGCGTGTGTTTCTAGGTCGTGGCGTCG 1909	gg Qy
1792 GGAACGGCGGTGGCGGTGCGGCGTCTCGGTGCGCTCTGGCCGCTCAGCCGTGGCGG 1849	g Qy
1733 CCGCCCCGTGGACGCGGGTTCCCGT-GGCGTTCCCGCGGCCAGGCATCAGCAATCTATCAG 1791	B 8
1674 ACCCGGCAGGCGCCCGGGAAACTCCGCCCCCGGCCGGCCG	og Qy
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554 TGADATCAAGTAAGACTTCCTCGGACAGAACAATCAAAGGGGTGCGCCGGGGTCCTCCAA 161	. B &
TTG 155	Db Qy
149 600	oy Oy
1374 TTTAATTAGCCAGTTGTGGTGAGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGTGGG 1433	Db dy
137 612	D S
1254 GGTAGCTCATGCCCCTGATCCCAGCACTTCGGGAGGCTGAGGGCGTGAAGATCACTTGTAG 1313	B &
1194 TTCCCGGACCCAGCTGCCTCAGGGGAGAGAGACACACTTAAGAGTTTGGGGCCGGCGT 1253	B 8
1134 CTAGCACTGCAGGGTCCACCTCGGGCCAATCAATTATTTCCCGAGGCGGGGGCCTGCGC 1193	B &
1074 TTTGTCCCGACGGTCTGTGGACGGCAGGTTAGAACGCCCGTCCAAGCCAGGAGGGTGGAC 1133	D Q

RESULT 7
US-10-311-455-44/c
; Sequence 44, Application US/10311455
; Publication No. US20030143606A1

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GENERAL INFORMATION:

APPLICANT: DIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases As:
TITLE OF INVENTION: Diagnosis of Diseases As:
TITLE OF INVENTION: Cytosine methylation
FILE REFERENCE: 5013.1014
CURRENT APPLICATION NUMBER: US/10/311,455
CURRENT FILING DATE: 2002-12-16
FRIOR APPLICATION NUMBER: PCT/EP01/07537
PRIOR FILING DATE: 2001-07-02
FRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 2424
LENGTH: 10619
TYPE: DNA
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Matches 1600;
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Best Local Similarity 76.8%;
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FEATURE:
OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
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                                            GAGTTTTATTCATCTGCTTATTTTTCTTATCCTCTATACCAGAATTGAATATTTTCAAAT
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    AAACCAATTTTAATAAACTATATTTTGAAGTATAGTTCTATATTAAACAACAAGATCTAG 720
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                                                          ANANCTACTANCTCNACCCANACCCCGCCTCGACCCCCA-AACAACGACCGCANAACTCC
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CURRENT APPLICATION NUMBER: US/10/240,453
CURRENT FILING DATE: 2002-10-02
PRIOR APPLICATION NUMBER: PCT/EP01/03973
PRIOR FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR FILING DATE: 2000-04-07
PRIOR PRILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR PRILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR PRILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 350
SEQ ID NO 2
SEQ ID NO 2
LENGTH: 10619
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US-10-240-453-2/c
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Publication No. US20030148326A1
GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: PIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA
TITLE OF INVENTION: Transcription
TITLE OF INVENTION: by Means of Assessing the Methylation Status
TITLE OF INVENTION: With DNA TRANSCription
FILE REFERENCE: 5013.1009
FILE REFERENCE: 5013.1009
                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 1600; Conserv
                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                 FEATURE:
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                 AAATATC--TTTTTTTTTTTTTAAAACAAAATCACACTATCACCCAAACTAAAATCCAA
                                                                                 TTCTCTAAATAAAACTTTTATTTATTTCCATCATATTTTCCACTTATTCTATTTACCTTCA
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                                                                                                                                                  TTTAAAAATAATATTATCAACCCAAAAATAACATAATCATACCTTTTAACTTAATCA
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                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                  TTTGTCCCGACGGTCTGTGGACGGCAGGTTAGAACGCCCGTCCAAGCCAGGAGGGTGGAC 1133
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Sequence 2, Application US/10240589C
Publication No. US20040076956A1
GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: PIEDENBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Diagnosis of Diseases Associated wit
TITLE OF INVENTION: DNA repair
FILE REFERENCE: 5013.1008
CURRENT APPLICATION NUMBER: US/10/240,589C
CURRENT FILING DATE: 2003-09-02
PRIOR APPLICATION NUMBER: PT/EP01/03972
PRIOR FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR FILING DATE: 2000-04-06
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PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 148
SEQ ID NO 2
LENGTH: 10619
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity
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                                                                    GCCAGGTGCAGTGGCTCATGCCTGTAATCCCCAGCAATTTTGGGAAGTCGAGGTGGGAGGAT 780
                                                                                                                        AAACCAATTTTAATAAACTATATTTTGAAGTATAGTTCTATATTAAACAACAACAACTATGT
                                                                                                                                                                    АДАЛСАСТСАТАТТАСЛАТСТТТАЛЛАТ-ДАЛЛАЛЛАЛЛАТАСАТАЛЛАТТАЛЛАЛЛА 6848
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CTAPATAAACGCACGCGAAACGACGAAACGACAA---CGTATTTCTAAATCGTAACGTCG
                      CTGGGTGAGCGCAGGCGGCGAGGCGGCAAGCGTGTGTTTCTAGGTCGTGGCGTCG
                                                                       AAAACGACGATAACCGATACGACGTATTCGATAACGACTCTAACCGCTCAAAAACTACGA
                                                                                                GGAACGGCGGTGGCCGCTGCTGCCGCTCTGGCCGCTCAGCCCTGGCCG
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CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR APPLICATION NUMBER: US 60/199,676
PRIOR FILING DATE: 2000-04-20
PRIOR PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR PILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
PRIOR APPLICATION NUMBER: US 60/146,002
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US-10-027-632-154183
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Publication No. US20020198371A1
GENERAL INFORMATION:
APPLICANT: Wang, David G.
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
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Best Local Similarity 96.6
Matches 822; Conservative
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TCAGTCTCAAAAATAAAAGGGGGAGGGGTTGGGGGGTAAAATTAGTTGTGAAATCAAGTAA
                                                                                                              TCAGGAGTTCCAGACTGCAGTGAGCCATGATGGCGGCACTGCACTCCAGCGCGCGGTGAGAC 1506
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Pred. No. 5.3e-161;
1; Mismatches 12;
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APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/193,632
CURRENT APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-00-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/193,636
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-09-28
PRIOR PR
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US-10-027-632-154183
; Sequence 154183, Application US/10027632
; Publication No. US20030204075A9
; GENERAL INFORMATION:
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 Query Match
                                                       LENGTH: 844
TYPE: DNA
ORGANISM: Human
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Sequence 107, Application US/09960253
Patent No. US20020123619A1
GENERAL INFORMATION:
APPLICANT: Benson, Darin R.
APPLICANT: Mohamath, Raodoh
APPLICANT: Lodes, Michael J.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
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Best Local Similarity 96.6 Matches 822; Conservative
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                                               CTCCGGATGGC 2049
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FILE REFERENCE: 210121.556
CURRENT APPLICATION NUMBER: US/09/960,253
CURRENT FILING DATE: 2001-09-20
NUMBER OF SEQ ID NOS: 187
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 107
LENGTH: 665
TYPE: DNA
ORGANISM: Homo sapiens
US-09-960-253-107
                                                                                                                       NUMBER OF SEQ ID NOS: 341
SOFTWARE: Corixa Invention Disclosure Database SEQ ID NO 300
LENGTH: 3859
TYPE: DNA
ORGANISM: Homo sapiens
US-09-864-864-300
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APPLICANT: Mitcham, Jenni
APPLICANT: Mitcham, Jenni
APPLICANT: Billon, Davin
APPLICANT: Secrist, Heath
APPLICANT: Lodge, Michael
APPLICANT: Algate, Paul A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. US20020102679A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 281; Conserv
                                                                                   Query Match 12.5%;
Best Local Similarity 97.2%;
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                                                                    Matches 277;
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APPLICANT:
APPLICANT:
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APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.523
CURRENT APPLICATION NUMBER: US/09/864,864
CURRENT APPLICATION DATE: 2001-05-23
                   1782 AATCTATCAGGGAACGGCGGTGGCGGTGCGGCGTTTCGGTGCGCTCTGGCCGCTCAGC 1841
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Fling, Steve P.
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Harlocker, Susan I
Dillon, Davin C.
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                                                                  Conservative
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                                                                 Score 261.4;
Pred. No. 3.8e
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Pred. No. 1.5e-54;
0; Mismatches 6
                                                             3.8e-53;
6;
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PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR PELICATION NUMBER: 60/323,580
PRIOR FILING DATE: 2001-09-19
NUMBER OF SEQ ID NOS: 363
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 3859
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APPLICANT: John MONAHAN
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                                                                             Matches 277;
                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification, TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer FILE REFERENCE: MRI-030
CURRENT APPLICATION NUMBER: US/10/097,340
CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: 60/276,025
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR APPLICATION NUMBER: 60/325,149
PRIOR APPLICATION NUMBER: 60/3276,026
PRIOR FILING DATE: 2001-03-14
PRIOR FILING DATE: 2001-08-16
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                                                                                                   Local Similarity
                     1782 AATCTATCAGGGAACGGCGGTGGCCGGTGCGGCGTTTCGGTGCGCTCTGGCCGCTCAGC 1841
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Shubhangi KAMATKAR
Steve G. KOVATS
Rachel E. MEYERS
Michael MORRISEY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gordon B. MILLS
Robert C. BAST, Jr.
Karen LU
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Karen GLATT
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                                                                                Conservative
                                                                                                12.5%;
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                                                                           Score 261.4; DB 14;
Pred. No. 3.8e-53;
0; Mismatches 6;
                                                                                                                      DB 14;
                                                                                Indels
                                                                                                                   Length 3859;
                                                                             <u>ب</u>
                                                                             Gaps
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SOPTWARE: PATENTIN Version 3.1
SEQ ID NO 3
LENGTH: 3859
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (160)..(3204)
OTHER INFORMATION:
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US-10-163-587A-3
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publication No. US20030096263A1
GENERAL INFORMATION:
APPLICANT: Oliveira, Marcos
TITLE OF INVENTION: SELECTIVE PARP-1 TARGETING FOR DESIGNING CHEMO/RADIO SENSITIZING
FILE REFERENCE: 50229-306
CURRENT APPLICATION NUMBER: US/10/163,587A
CURRENT FILING DATE: 2003-01-10
PRIOR APPLICATION NUMBER: 60/296,110
PRIOR APPLICATION NUMBER: 60/296,110
PRIOR FILING DATE: 2001-06-07
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 12.5%; Score 261.4; DB 14; Length 3859; Best Local Similarity 97.2%; Pred. No. 3.8e-53; Matches 277; Conservative 0; Mismatches 6; Indels 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1782 AATCTATCAGGGAACGGCGGTGGCCGGTGCGGCGTTCGGTGCGCTCTGGCCGCTCAGC 1841
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                                                                                                                    2022 GCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAGGTGC 2066
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Result
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence:
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         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Match
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1: /ggn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
         GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.
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                          US-08-222-177A-445
US-08-222-177A-146
US-08-222-177A-143
US-08-222-177A-143
US-08-222-177A-451
US-08-222-177A-451
US-08-222-177A-238
US-08-222-177A-238
US-08-222-177A-215
US-08-222-177A-215
US-08-222-177A-215
US-08-222-177A-216
US-08-222-177A-216
US-08-222-177A-376
US-08-222-177A-170
US-08-222-177A-170
US-08-222-177A-110
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Sequence 445, App
Sequence 143, App
Sequence 23, Appl
Sequence 23, Appl
Sequence 23, Appl
Sequence 23, Appl
Sequence 25, Appl
Sequence 80, Appl
Sequence 80, Appl
Sequence 218, Appl
Sequence 218, Appl
Sequence 271, Appl
Sequence 215, Appl
Sequence 217, Appl
Sequence 217, Appl
Sequence 218, Appl
Sequence 210, Appl
Sequence 210, Appl
Sequence 210, Appl
Sequence 110, Appl
Seq
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Query Match 100.0%; Score 24; DB 1; Length 24; Best Local Similarity 100.0%; Pred. No. 0.27; Matches 24; Conservative 0; Mismatches 0; Indels	INFORMATION FOR SEQ ID NO: 445: SEQUENCE CHARACTERISTICS: LENGTH: 24 base pairs TYPE: nucleic acid TYPE: nucleic acid TRANDEDNESS: double TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) US-08-222-177A-445	TE AT	APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 435 PRIOR APPLICATION DATA, APPLICATION NUMBER: FILING DATE: 21 APPLICATION DATA	; CITY: Maddson ; STATE: Wisconsin ; COUNTRY: USA ZIP: 53717-1914 ; COMPUTER READABLE FORM: ; MEDIUM TYPE: Floppy disk ; COMPUTER: IBM PC compatible ; COMPUTER: IBM PC compatible ; OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.25 ; CURRENT APPLICATION DATA:	GENERAL IN APPLICAN TITLE OF TITLE OF NUMBER C CORRESPC ADDRES STREET	ALIGNMENTS RESULT 1 US-08-222-177A-445 ; Sequence 445, Application US/08222177A ; Patent No. 5582979	28 24 100.0 36 1 US-08-222-177A-158 Sequence of the control of the
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Gaps					USING		358 App 140 App 68, App 68, App 198 App 137, App 137, App 137, App 152, App 175, App 176, App 177, App 177, App 178, App 179, App 170, App 170, App 170, App 170, App
0;					SAME		200000000000000000000000000000000000000

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US-08-455-627-23
, Sequence 23, Application US/08455627
; Patent No. 5571677
; GENERAL INFORMATION: APPLICANT: Sergei M. Gryaznov
; TITLE OF INVENTION: Convergent Synthesis of Branched and Multiply
; TITLE OF INVENTION: Connected Macromolecular Structures
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US-08-222-177A-146
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GENERAL INFORMATION:
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CLASSIFICATION: 435
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US 07/341,562
APPLICATION NUMBER: US 07/341,562
APPLICATION INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 09.492
REFERENCE/DOCKET NUMBER: 09.65.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                        NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: DI
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ADDRESSEE: Cooley Godward LLP STREET: Five Palo Alto Square, 3000 El Camino Real CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 8000 H
                                                                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                           24;
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                                                                                                                                                                                                                                                                                                                                                                                           h 100.0%; Score 24; DB 1 Similarity 100.0%; Pred. No. 0.27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25 base pairs
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WEDER, James L.
VENTION: LENGTH POLYMORPHISMS IN
VENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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US-08-222-177A-143; Sequence 143, Application US/08222177A; Patent No. 5582979
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Best Local S
Matches 24
                        APPLICATION NUMBER: US 07/341
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 0986
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 415-857-0663
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 27 nucleotides
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-BOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: LENGTH POLYMORPHISMS IN TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: LY
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-843-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/455,627 FILING DATE: 31-MAY-1995 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION: NAME: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Weber, James L.
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STRANDEDNESS: sing
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REGISTRATION NUMBER: 35,966
TELEFAX:
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8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
    (608) 831-2106
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SAME

INFORMATION

SEQUENCE CHARACTERISTICS

FOR SEQ ID NO:

143:

ENGTH:

27 base pairs

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US-08-689-856-23
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                                                                                                                                                                                            TELEFAX: 415-0... 23
INFORMATION FOR SEQ ID NO: 23
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
1.ENGTH: 27 nucleotides
                                                              Best Loc
Matches
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GENERAL INFORMATION:
                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/.
FILING DATE: 31-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Nakamura, Jackie N.
REGISTRATION NUMBER: 35,96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: LY TELECOMMUNICATION INFORMATION: TELEPHONE: 415-843-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Convergent Synthesis of Branched and Multiply TITLE OF INVENTION: Connected Macromolecular Structures NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooley Godward LLP
STREET: Five Palo Alto Square, 3000 El Camino Real
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMMEDIATE SOURCE:
CLONE: mfd31rs
                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                                                                            Local
                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ch 100.0%; Score 24; DB 1 l Similarity 100.0%; Pred. No. 0.27; 24; Conservative 0; Mismatches
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                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Palo Alto
                             CACACACACACACACACACACACA
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                                                              Conservative
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                                                            100.0%; Score 24;
100.0%; Pred. No.
tive 0; Mismatch
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                                                              Mismatches
                                                                            DB 2; Length 27; 0.27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-222-177A-451
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CURRENT APPLICATION NUMBER: US/08/787,321A
CURRENT FILING DATE: 1997-01-03
EARLIER APPLICATION NUMBER: US PROV 60/009,918
EARLIER FILING DATE: 1996-01-12
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 451, Application US/08222177A Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Patent No. 6180777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                           APPLICATION NUMBER: US 07/341
ETLING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 0986
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION: APPLICANT: Weber,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: HORN, Thomas
TITLE OF INVENTION: SYNTHESIS OF BRANCHED NUCLEIC ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
STRANDEDNESS:
                LENGTH: 28 base pairs TYPE: nucleic acid
                                                                                                               TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Madison
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                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 24; DB 3; Length 27; 100.0%; Pred. No. 0.27;
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; TOPOLOGY: linear; MOLECULE TYPE: DNA (genomic) US-08-222-177A-451
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                                                                                                                                                                                                                                                                                                                                                                             US-08-455-627-25
Sequence 80, Application US/08222177A
Patent No. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
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Patent No.
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Best Local
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Sergei M. Gryaznov
TITLE OF INVENTION: Convergent Synthesis of Branched and Multiply
TITLE OF INVENTION: Connected Macromolecular Structures
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooley Godward LLP
STREET: Five Palo Alto Square, 3000 El Camino Real
                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: 1
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 29 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: NAKAMULTA, Jackie N.
REGISTRATION NUMBER: 35,966
REFERENCE/DOCKET NUMBER: LYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94306-2155
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-843-5000
TELEFAX: 415-857-0663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/455,627
FILING DATE: 31-MAY-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25, Application US/08455627
5. 5571677
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100.0%; Pred. No.
tive 0; Mismatch
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0.27;
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US-08-222-177A-238
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GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341
PILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: DNI IMMEDIATE SOURCE: CLONE: mfdl0rs
                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                          COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401 CITY: Madison STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 53717-1914
                                                                                                                                                                                                                COUNTRY: USA
ZIP: 53717-1914
                                                            FILING DATE:
                                                                                                                                                                                                                                                                           CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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                                                                                                                                                                                                                                                         Wisconsin
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                                                                                                                                                                                                                                                                                             B: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                 Weber, James L.

VENTION: LENGTH POLYMORPHISMS IN

VENTION: (dC-dA)n.(dG-dT)n SEQUE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 24; DB 1; Length 29; 100.0%; Pred. No. 0.27;
                                                                                                                                                                                                                                                                                                                                                       460
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    US 07/341,562
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REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100

FILING DATE: 21-APR-1989 ATTORNEY/AGENT INFORMATION: NAME: Sara, Charles S.

REGISTRATION NUMBER:

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RESULT 11
US-08-689-856-25
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US-08-689-856-25
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Sequence 25, App.
Sequence 25, App.
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Best Local
                                                                                 INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION: APPLICANT: Sergei
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                              ZIP: 94306-2155
COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/689,856
FILING DATE:
                                                                                                                                                        NAME: Nakamura, Jackie N.
REGISTRATION NUMBER: 35,966
REFERENCE/DOCKET NUMBER: LY:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-843-5000
                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/455,627
FILING DATE: 31-MAY-1995
ATTORNEY/AGENT INFORMATION:
ANALY MACHINERY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooley Godward LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Convergent Synthesis of Branched and Multiply TITLE OF INVENTION: Connected Macromolecular Structures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                STRANDEDNESS: single TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Five Palo Alto Square, 3000 El Camino Real
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE:
                                                                                                                                             TELEFAX: 415-857-0663
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                                                                   nucleic acid
               TYPE:
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100.0%; Pred. No. 0.27;
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                                                                                                     US-08-222-177A-215
                                                                                                                        RESULT 13
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                                                                      Sequence 215, Application US/08222177A Patent No. 5582979
                                                                                                                                                                                                                                           Matches
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Best Local S
                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dc-da)n. (dg-dt)n sequences and methods of using same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION: APPLICANT: Weber,
                                                                                                                                                                                                                                                                                                                           IMMEDIATE SOURCE:
CLONE: mfd118rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Madison
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH: 30 base pairs
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                                                                                                                                                                                                                                                           100.0%;
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                                                                                                                                                                                                                                                           Score 24; I
Pred. No. 0.
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                                                                                                                                                                                                                                                             DB 1;
0.27;
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CORRESPONDENCE ADDRESS: NUMBER OF SEQUENCES:

460

ADDRESSEE:

E: DeWitt Ross & Stevens, S.C. 8000 Excelsior Drive, Suite 401

CITY: Madison

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RESULT 14
US-08-222-177A-235
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Patent No. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
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FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/341,562
FILING DATE: 21-APR-1989
AITORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
NAME: Sara, Charles S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 31 base pairs
                                 COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: (d
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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TOPOLOGY: linear
MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                            STREET:
                     FILING DATE:
                                                                                                                                                                                COUNTRY: USA
ZIP: 53717-1914
                                                                                                                                                                                                                                             CITY: Madison
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CLASSIFICATION:
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8000 Excelsior Drive, Suite 401
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Best Local Similarity
Matches 24; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dc-da)n. (dg-dT)n SEQUENCES AND METHODS OF USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 31 base pairs
                                                   INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
PILING DATE: 21-APR-1989
ATTORNEY_AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 09865.601
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                 ZIP: 53717-1914

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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CLONE: mfd64rs
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NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
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           LENGTH: 31 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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STRANDEDNESS:
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                                                                                                           (808)
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                                                                          271:
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Perfect score:
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Gapop 10.0 , Gapext 1.0
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12:
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1 cacacacacacacacacacacaca 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Published Applications NA:*

1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/2/pubpna/PCT NEW PUB.seq:*
2: /cgn2_6/ptodata/2/pubpna/US06_NEW PUB.seq:*
3: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
4: /cgn2_6/ptodata/2/pubpna/US07_NEW PUB.seq:*
5: /cgn2_6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
6: /cgn2_6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
Cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US09E_PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US11NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US11NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US11NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US10F_PUBCOMB.seq:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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(without alignments)
6202.889 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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24	24	24	24	24	24	24	Score
100.0	100.0	100.0	100.0	100.0	100.0	100.0	Query Match Length DB ID
24	24	24	24	24	24	24	Length
17				11	10	9	8
US-10-314-578-1068	US-10-017-995-1068	US-10-112-653-1012	US-09-909-317-6	US-09-776-479-1068	US-09-776-479-1068	US-09-735-363A-21	DB ID
Sequence 1068, Ap	Sequence 1068, Ap	Sequence 1012, Ap	Sequence 6, Appli	Sequence 1068, Ap	Sequence 1068, Ap	Sequence 21, Appl	Description

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Sequence 775, App Sequence 29, Appl Sequence 28, Appl Sequence 114, App Sequence 234, App Sequence 236, App	Sequence 26, Sequence 27, Sequence 27, Sequence 28, Sequence 28, Sequence 33, Sequence 514, equence 571, eque	Sequence 25, Sequence 24, Sequence 24, Sequence 24, Sequence 24, Sequence 24, Sequence 24, Sequence 24,	Sequence 27, Appl Sequence 21, Appl Sequence 10, Appl Sequence 22, Appl Sequence 23, Appl Sequence 74, Appl Sequence 678, App	equence 1068 quence 1, Ap quence 5, Ap quence 66, 1 quence 770, equence 2, 1 equence 93,

ALIGNMENTS

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RESULT 1
US-09-735-363A-21/c
; Sequence 21, Application US/09735363A
; Patent No. US20010041681A1
; GENERAL IMPORMATION:
APPLICANT: Filion, Mario
; PILE REFERENCE: 0281-0181
; CURRENT APPLICATION NUMBER: 60/170,325
; PRIOR APPLICATION NUMBER: 60/170,325
; PRIOR FILING DATE: 1999-12-13
; PRIOR FILING DATE: 1999-12-13
; PRIOR FILING DATE: 2000-08-29
; NUMBER OF SEQ ID NOS: 87
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 21
; LENGTH: 24
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER IMPORMATION: Synthetic Oligonucleotide
US-09-735-363A-21

Query Match
Best Local Similarity 100.0%; Score 24; DB 9; Length 24;
Best Local Similarity 100.0%; Pred. No. 0.47;
Matches 24; Conservative 0; Mismatches 0; Indels 0; Gaps

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24

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APPLICANT: Bratzler, Robert L.
APPLICANT: Petersen, Deanna M.
APPLICANT: Pouron, Yves
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR APPLICANTS
PRIOR OF SECULO NUMBER: US 60/179,991
RESULT 4
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US-09-776-479-1068/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 1068 LENGTH: 24
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Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bratzler, Robert L.
APPLICANT: Petersen, Deanna M.
APPLICANT: Petersen, Deanna M.
APPLICANT: Fouron, Yves
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR APPLICATION NUMBER: US 60/179,991
RIOR FILING DATE: 2000-02-03
NUMBER OF SEG ID NOS: 1093
                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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100.0%; Pred. No.
tive 0; Mismatch
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APPLICANT: Krieg, Arthur M.
APPLICANT: Berg, Daniel J.
APPLICANT: Berg, Daniel J.
APPLICANT: Berg, Daniel J.
APPLICANT: BOR INMUNOSTIMULATORY NUCLEIC ACID FOR
IITLE OF INVENTION: IRMUNOSTIMULATORY NUCLEIC INFLAMMATORY DISEASES
FILE REFERENCE: C01039/70060(AWS)
CURRENT APPLICATION NUMBER: US/10/112,653
CURRENT FILING DATE: 2002-03-29
PRIOR APPLICATION NUMBER: US 60/279,642
PRIOR APPLICATION NUMBER: US 60/279,642
PRIOR APPLICATION NUMBER: US 60/279,642
PRIOR FILING DATE: 2001-03-29
NUMBER OF SEQ ID NOS: 1040
SOPTWARE: FRACESEQ for Windows Version 3.0
SEQ ID NO 1012
LENGTH: 24
TYPE: DNA
TYPE: DNA
TYPE: DNA
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APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
TITLE OF INVENTION: Genetic Marker Test for
FILE REFERENCE: 18810-82152
CURRENT APPLICATION NUMBER: US/09/909,317
CURRENT FILING DATE: 2001-07-18
PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 7
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GENERAL INFORMATION:
APPLICANT: Bratzler, Robert L.
TITLE OF INVENTION: Inhibition of Angiogenesis
FILE REFERENCE: C1037/7025 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/10/017,995
CURRENT FILING DATE: 2001-12-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1012, Application US/10112653 Publication No. US20030050268A1 GENERAL INFORMATION:
                                                                                                                               Sequence 1068, Application US/10017995 Publication No. US20030055014A1
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Best Local Similarity
Matches 24; Conserv
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Best Local
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Publication No. US20040152075A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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Pred. No. 0.47;
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Sequence 1068, Application US/10831778

| Publication No. US20040235774A1
| GENERAL INFORMATION:
| APPLICANT: Bratzler, Robert L.
| APPLICANT: Bratzler, Deanna M.
| APPLICANT: Fouron, Yves
| TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
| TITLE OF INVENTION: Treatment of Asthma and Allergy
| FILE REFERENCE: C1037/7013 (HCL/MAT)
| CURRENT APPLICATION NUMBER: US/10/831,778
| CURRENT FILING DATE: 2004-04-23
| PRIOR APPLICATION NUMBER: US 60/179,991
| PRIOR FILING DATE: 2000-02-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Krieg, Arthur M.
APPLICANT: Schetter, Christian
APPLICANT: Vollmer, Jorg
TITLE OF INVENTION: Immunostimulatory Nucleic Acids
FILE REFERENCE: C1039/7035 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/10/314,578
CURRENT FILING DATE: 2002-12-09
PRIOR APPLICATION NUMBER: US 60/156,113
PRIOR FILING DATE: 1999-09-25
PRIOR APPLICATION NUMBER: US 60/156,135
PRIOR FILING DATE: 1999-09-27
PRIOR APPLICATION NUMBER: US 60/227,436
PRIOR APPLICATION NUMBER: US 60/227,436
PRIOR FILING DATE: 2000-08-23
NUMBER OF SEO ID NOS: 1145
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1068
ENOTH: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-017-995-1068
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SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1068
LENGTH: 24
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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Pred. No. 0.47;
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GENERAL INFORMATION:

APPLICANT: Filion, Mario

APPLICANT: Fhilip, Nigel

ITITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides

FILE REFERENCE: 02811-0181

CURRENT APPLICATION NUMBER: US/09/735,363A

CURRENT FILING DATE: 2000-12-12

PRIOR APPLICATION NUMBER: 60/12-0,325

PRIOR APPLICATION NUMBER: 60/228,925

PRIOR APPLICATION NUMBER: 60/228,925

PRIOR APPLICATION NUMBER: 00/28,925

PRIOR FILING DATE: 2000-08-29

NUMBER OF SEQ ID NOS: 87

SOFTWARE: PatentIn version 3.0

SEQ ID NO 1

LENGTH: 27
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              SOFTWARE: Pat
SEQ ID NO 5
SEQ TO NO 5
                                                                                                                                                                                                                                                                                     Sequence 5, Application US/09735363A Patent No. US20010041681A1 GENERAL INFORMATION:
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Best Local Similarity
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SOFTWARE: FREESEQ for Windows Version 3.0
SEQ ID NO 1068
LENGTH: 24
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                                                                                                                                                                  APPLICANT: Filion, Mario
APPLICANT: Phillip, Nigel
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
FILE REFERENCE: 02811-0181
CURRENT APPLICATION NUMBER: US/09/735,363A
CURRENT FILING DATE: 2000-12-12
                                                                                                            PRIOR APPLICATION NUMBER: 60/170,325
PRIOR FILING DATE: 1999-12-13
PRIOR APPLICATION NUMBER: 60/228,925
                                                                          PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
TYPE: DNA
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                                                      PatentIn version 3.0
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100.0%; Pred. No. 0.47;
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CURRENT APPLICATION NUMBER: US/09/735,363A
CURRENT FILING DATE: 2000-12-12
PRIOR APPLICATION NUMBER: 60/170,325
PRIOR FILING DATE: 1999-12-13
PRIOR APPLICATION NUMBER: 60/228,925
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 87
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 66
ENOTH: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Synthetic Oligonucleotide US-09-735-363A-5
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US-09-735-363A-66
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                                                                                                                                                                                                                                                                                                                                                                             Sequence 770, Application US/09263959 Patent No. US20020150891A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 66, Application US/09735363A Patent No. US20010041681A1 GENERAL INFORMATION:
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Best Local &
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Best Local Similarity
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APPLICANT: Phillip, Nigel
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
FILE REFERENCE: 02811-0181
                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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           CURRENT APPLICATION DATA: US
                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Flappy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                             APPLICANT: ROWEN, Lee

APPLICANT: KOOD, Ben F.

TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
NUMBER OF SEQUENCES: 1279
                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                        ZIP: 98104-7092
                                                                                                                                                                               CITY: Seattle
STATE: Washing
                                                                                                                                                                                                                STREET:
                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                               ADDRESSEE:
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1: US
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05-MAR-1999
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           US/09/263,959
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                                                   Version #1.25
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0.47;
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APPLICANT: Gray, Gary S.
TITLE OF INVENTION: POLYMORPHIC ELEMENTS IN THE TITLE OF INVENTION: COSTIMULATORY RECEPTOR LOFILE REFERENCE: GNN-5343CP2
CURRENT APPLICATION NUMBER: US/10/085,906
CURRENT FILING DATE: 2002-02-27
PRIOR APPLICATION NUMBER: US 60/126,215
PRIOR APPLICATION NUMBER: US 09/534,061
                                                                                                                                                                                                                                                                       US-10-085-906-93/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Phillips, Nigel C.
APPLICANT: Pillon, Mario C.
TITLE OF INVENTION: Hyaluronic Acid in the Treatment of Cancer FILE REFERENCE: 02811-0211 (42368-274915)
CURRENT APPLICATION NUMBER: US/10/168,327
CURRENT FILING DATE: 2002-10-07
PRIOR APPLICATION NUMBER: PCT/CA00/01562
PRIOR FILING DATE: 2000-12-28
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
                                                                                                                                                     Sequence 93, Application US/10085906
Publication No. US20030054371A1
GENERAL INFORMATION:
APPLICANT: Ying, Vincent
APPLICANT: Wu, Paul
APPLICANT: Gray, Gary S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2, App. Publication No.
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 27 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Synthetic oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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REGISTRATION NUMBER: 3:
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Local Similarity 100.0%; Pred. No. 0.47;
es 24; Conservative 0; Mismatches
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; PRIOR FILING DATE: 2000-03-24
; PRIOR APPLICATION NUMBER: PCT/US00/07938
; PRIOR FILING DATE: 2000-03-24
; NUMBER OF SEQ ID NOS: 545
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 93
; LENGTH: 30
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-085-906-93
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Search completed: September Job time: 27.3388 secs
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US-10-085-906-27/c
                                                                             Query Match 100.0%; Score 24; DB 14; Length 31; Best Local Similarity 100.0%; Pred. No. 0.47; Matches 24; Conservative 0; Mismatches 0; Indels
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Scoring table:
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1: /cgn2_6/ptcdata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptcdata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptcdata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptcdata/1/ina/6B_COMB.seq:*

5: /cgn2_6/ptcdata/1/ina/PCTUS_COMB.seq:*

6: /cgn2_6/ptcdata/1/ina/backfiles1.seq:*
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Maximum Match 100%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
            GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd
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  US-08-222-177A-140
US-08-222-177A-397
US-08-222-177A-137
US-08-222-177A-119
US-08-222-177A-403
US-08-222-177A-403
US-08-222-177A-23
US-08-222-177A-23
US-08-222-177A-23
US-08-222-177A-23
US-08-222-177A-29
US-08-222-177A-195
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US-08-222-177A-346
US-08-222-177A-36
US-08-222-177A-39
US-08-222-177A-36
US-08-222-177A-36
US-08-222-177A-36
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Sequence 140, App Sequence 137, App Sequence 137, App Sequence 400, App Sequence 400, App Sequence 403, App Sequence 183, App Sequence 232, App Sequence 232, App Sequence 239, App Sequence 195, App Sequence 195, App Sequence 101, App Sequence 1019, App Sequence 1019, App Sequence 101, App Sequence 210, App Sequence 210, App Sequence 310, App Sequence 33, App1 Sequence 33, App1 Sequence 22, App Sequence 33, App1 Sequence 21, App Sequence 21, App Sequence 21, App Sequence 31, App1 Sequence 21, App Sequence 361, A
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US-08-222-177A-359 US-08-222-177A-367 US-08-222-177A-89 US-08-222-177A-317 US-08-222-177A-317 US-08-222-177A-337 US-08-222-177A-337 US-08-222-177A-391 US-08-222-177A-391 US-08-222-177A-397 US-08-222-177A-307 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-394 US-08-222-177A-395 US-08-222-177A-394 US-08-222-177A-395 US-08-2

ALIGNMENTS

RESULT 1 US-08-222-177A-140

Sequence 140, Application US/08222177A Patent No. 5582979

US-08-222-177A-140 Query Match
Best Local Similarity
Matches 36; Conserv INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: GENERAL INFORMATION: APPLICANT: Weber, MOLECULE TYPE: DIMMEDIATE SOURCE: CLONE: mfd30rs NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 099
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/222,177A
APPLICATION NUMBER: US/08/222,177A PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION: APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH
TITLE OF INVENTION: (dC-dA COMPUTER READABLE FORM: NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS: STREET: o. STREET: Madison COUNTRY: ZIP: 5371 STRANDEDNESS: double TOPOLOGY: linear TYPE: nucleic acid TELEFAX: CLASSIFICATION: 435 FILING DATE: ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401 53717-1914 Wisconsin (608) 831-2106 USA DNA 100.0%; (genomic) LENGTH POLYMORPHISMS IN (dC-dA)n. (dG-dT)n SEQUENCES AND METHODS OF USING SAME 460 09865.601 Score 36; I Pred. No. 7. DB 1; I 7.8e-05; nes 0; Length 37;

Conservative

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Mismatches

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                                                                                                                        Sequence 397, Application US/08222177A Patent No. 5582979
GENERAL INFORMATION:
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                               Matches
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/341,562
PILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERNCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                 APPLICANT: Weber, James L.

TITLE OF INVENTION: LENGTH POLYMORPHISMS IN

TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES

NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n. (dG-dT)n SEQUENCES AND METHODS OF USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 38 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLONE: mfd50rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 53717-1914
ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (608) 831-2106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Madison
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                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                           Wisconsin
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8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
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DeWitt Ross & Stevens, S.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 36; DB 1; 100.0%; Pred. No. 7.8e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION: APPLICANT: Weber,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 38 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 0986
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
TELEPHONE: (608) 831-2106
                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
FILLING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                           ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLILATE: 21-Ar. - FILING DATE: 21-Ar. - ATTORNEY/AGENT INFORMATION: Sara, Charles S.
                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 8000 E. CITY: Madison STATE: Wiscons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 38 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                          CITY: Madison
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                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  137, Application US/08222177A
o. 5582979
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                                                                                                                                                                                                                                                                                                                                       E: DeWitt Ross & Stevens, S.C. 8000 Excelsior Drive, Suite 401
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
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Pred. No. 7.8e-05;
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REGISTRATION NUMBER:

30,492

09865.601

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US-08-222-177A-119
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US-08-222-177A-119
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                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/341,562

PILING DATE: 21-APR-1989

ATTORNEY/AGENT INFORMATION:

NAME: Sara, Charles S.

REGISTRATION NUMBER: 09.492

REFERENCE/DOCKET NUMBER: 09865.601

TELECOMMUNICATION INFORMATION:

TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATION:
COMPUTER: PATENT: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/222,177A
APPLICATION NUMBER: US/08/222,177A
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                        TOPOLOGY: line
MOLECULE TYPE: D
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: D
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 8000 P
                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                      FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 53717-19
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TOPOLOGY: linear
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                                                                                            nucleic acid
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               mfd23re
                                                                                                               40 base pairs
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8000 Excelsior Drive, Suite 401
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                                                              linear
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                                             DNA (genomic)
                                                                              double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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Pred. No. 7.8e-05;
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RESULT 7
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           Sequence 403, Application US/08222117A
PACENT NO. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n. (dG-dT)n SEQUENCES:
NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 400, Application US/08222177A Patent No. 5582979
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341
PILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 0986
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION: APPLICANT: Weber,
                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: 1: MOLECULE TYPE:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 8000 |
CITY: Madison
                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                    linear
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                                                                                                                                                                                                                                                                    100.0%; Score 36; DB 1; 100.0%; Pred. No. 7.8e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH POLYMORPHISMS IN (dC-dA)n. (dG-dT)n SEQUENCES AND METHODS OF USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          460
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Pred. No. 7.8e-05;
                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                         37
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                                                                                                                                                                                                                                                                                   Length 40;
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                            AND METHODS OF USING SAME
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ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401 CITY: Madison

Wisconsin

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RESULT 8
US-08-222-177A-74
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GENERAL INFORMATION:
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FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
PILING DATE: 21-APR-1999
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
APPLICATION: 435
EXIGN APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
APPLICATION NUMBER: 1989
                                                                                                         COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/222,177A
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                            CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX:
                                                                                                                                                                                                                                                                           STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CACACACACACACACACACACACACACACACACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (608) 831-2106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 36; 100.0%; Pred. No.
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; CLONE: mfd8rs
US-08-222-177A-74
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US-08-222-177A-183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 183, Applicat Patent No. 5582979
GENERAL INFORMATION:
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                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                              TOPOLOGY: line MOLECULE TYPE: DIMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Weber,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: double TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                       TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                   TELEFAX: (608) 831-2106
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                                                                                                                           LENGTH:
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5. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ch 100.0%; Score 36; DB 1; 1 Similarity 100.0%; Pred. No. 7.8e-05; 36; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     53717-1914
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                                                                                                                         41 base pairs
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8000 Excelsior Drive, Suite 401
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                                                                linear
                                                   DNA (genomic)
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US-08-222-177A-53
; Sequence 53, Application US/08222177A
; Patent No. 5582579
; Patent No. 5582579
; GENERAL INFORMATION:
; APPLICANT: Weber, James L.
; TITLE OF INVENTION: LENGTH POLYMO!
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                                                                                                                                                                                                                                                               ; IMMEDIATE SOURCE:
; CLONE: mfd63rs
US-08-222-177A-232
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US-08-222-177A-232
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, '
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                   TELEX:
[NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Madison
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                          ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Weber, James L.
VENTION: LENGTH POLYMORPHISMS IN
VENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                      (803)
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                                                                                                                                                                                                                                                                                                                             linear
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08) 831-2106
                                                                                                                                                                                                                                                                                                            DNA (genomic)
                                                                                                                                                                                                                                                                                                                                           double
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100.0%; Pr
 LENGTH POLYMORPHISMS IN
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Pred. No. 7.8e-05;
                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                 7.8e-05;
                                                                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                                                               Length 41;
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US-08-222-177A-229
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                                                                                                                                                                                                                                                                                                                        Patent No. 5562.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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MOLECULE TYPE:
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ZIP: 53717-1914
COMPUTER READABLE FORM:
COMPUTEN READABLE FORM:
TYPE: Floppy disk
                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: (CORRESPONDENCE ADDRESS:
                                                                                                                                                                  CITY: Madison
STATE: Wisconsin
                                                                                                                                 COUNTRY: USA
ZIP: 53717-1914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLONE: mfd1rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
                                                                                                                                                                                                                                                                                                                                                            229, Application US/08222177A
5. 5582979
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Similarity 100.0%;
36; Conservative 0
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                                                                                                                                                                                                       E: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (608) 831-2106
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                                                                                                                                                                                                                                                                                 (dC-dA)n. (dG-dT)n SEQUENCES
                                                                                                                                                                                                                                                                                                     LENGTH POLYMORPHISMS IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (\mathtt{dC-dA})\,n.\,(\mathtt{dG-dT})\,n SEQUENCES AND METHODS OF USING SAME 460
                                                                                                                                                                                                                                                                  460
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 36; DB 1;
Pred. No. 7.8e-05;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Version #1.25
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                                                                                                                                                                                                                                                                                 AND METHODS OF USING SAME
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 21-APR-1989

US 07/341,562

CLASSIFICATION:

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RESULT 13
US-08-222-177A-388
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; CLONE: mfd62rs
US-08-222-177A-229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 388, Application US/08222177A Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                 FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
APPLICATION NUMBER: US 07/341,562
APPLICATION UNFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 09.492
REFERENCE/DOCKET NUMBER: 09.65.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-da)n. (dG-dT)n SEQUENCES AND METHODS OF USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX:
INFORMATION FOR SEQ ID NO:
                TELEX:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 43 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 21-AFK-139
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                           ZIP: 53717-1914

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 8000 E:
CITY: Madison
STATE: Wiscons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (608) 831-2106
                                                                                                                  TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  36;
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                                                                               388
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US-08-222-177A-195
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US-08-222-177A-388
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                                                                                                                                    US-08-222-177A-195
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GENERAL INFORMATION:
APPLICANT: Weber,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100. Matches 36; Conservative
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                                                                   Matches
                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 0986
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: DNA
IMMEDIATE SOURCE:
CLONE: mfdl23rs
                                                                                                                                                TOPOLOGY: linea:
MOLECULE TYPE: DN
IMMEDIATE SOURCE:
CLONE: mfd49rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 53717-1914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Madison
STATE: Wiscons
                                                                                                                                                                                                                                   TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                          TELEX:
                                                                                                                                                                                                                                                                                                                          TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CACACACACACACACACACACACACACACACACA
                   36; Conservative
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                                                                                                                                                                                                    linear
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                                                                                                                                                                                    DNA (genomic)
                                                                                                                                                                                                                     double
                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 36; DB 1; 100.0%; Pred. No. 7.8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              US 07/341,562
                                                                   <u>.</u>
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                                                                                  Score 36; DB 1;
Pred. No. 7.8e-05;
                                                                   Mismatches
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RESULT 15 US-08-222-177A-241

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| Sequence 241, Application US/08222177A |
| Patent No. 558279 |
| GENERAL INFORMATION: | GC-ADT) NEQUENCES AND METHODS OF USING SAME |
| TITLE OF INVENTION: | GC-ADT) NEQUENCES AND METHODS OF USING SAME |
| TITLE OF INVENTION: | GC-ADT) NEQUENCES AND METHODS OF USING SAME |
| TITLE OF INVENTION: | GC-ADT) NEQUENCES AND METHODS OF USING SAME |
| TITLE OF INVENTION: | GC-ADT) NEQUENCES |
| APELICATION DATES: | 460 |
| CORRESPONDENCE ADDRESS: |
| ADDRESSEE: | DeWilt Rose & Stevens | S.C. |
| STREET: | 800 Excelsior Drive, Suite 401 |
| CITY: | Madison |
| STATE | Wisconsian |
| STATE | Wisconsian |
| COMPTRY: | USA |
| COMPTRY: | USA
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POSE BIONK (USPIO)

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Result
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Minimum DB
Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Searched:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OM nucleic - nucleic search, using sw model
                                                                       O
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                                                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                    Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   seq length: 0
seq length: 2000000000
      \begin{smallmatrix} \mathbf{G} & \mathbf{G} & \mathbf{G} & \mathbf{G} & \mathbf{G} \\ \mathbf{G} & \mathbf{G} & \mathbf{G} & \mathbf{G} & \mathbf{G} \\ \mathbf{G} & \mathbf{G} & \mathbf{G} & \mathbf{G} & \mathbf{G} \\ \end{smallmatrix}
                                                                                                                                                                                 Query
Match
                           100.0
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Gapop 10.0 , Gapext 1.0
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36
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                                                                                                                                                                                                                                                                                                                                                                        / Ggn2_6/ptodata/2/pubpna/USOO_RUBLEDB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOO_RUBLEDB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOB_PUBCOMB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOB_PUBCOMB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOB_PUBCOMB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOB_PUBCOMB.Seq:*
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/ Ggn2_6/ptodata/2/pubpna/USOSD_RUBLEDGMB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOSD_RUBLEDGMB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOSD_RUBLEDGMB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOSD_PUBCOMB.Seq:*
/ Ggn2_6/ptodata/2/pubpna/USOSD_NEW_PUB.Seq:*
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/cgn2_6/ptodata/2/pubpna/PCT_MEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USO6_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/USO6_PUBCOMB.seq:*
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11
  0 US-09-852-903C-23

US-09-909-317-7

0 US-09-852-903C-24

US-09-263-959-678

0 US-09-852-903C-25

9 US-10-661-088-24

9 US-10-661-097-24
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Sequence 23, Appl
Sequence 7, Appli
Sequence 24, Appli
Sequence 678, App
Sequence 25, Appli
Sequence 24, Appli
Sequence 24, Appli
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19 10 10 10 10 10 10 10 10 118 18	10 10 10 10 10		
US-10-407-818-15 US-09-852-903C-34 US-09-971-253-29 US-09-852-903C-35 US-09-852-903C-36 US-09-852-903C-37 US-09-854-867-450 US-10-786-970A-450 US-10-035-833A-5812 US-10-035-833A-6881 US-10-674-124A-25942	-10-407-814 -10-484-784 -09-971-353 -09-814-353 -09-852-903 -09-852-903 -09-852-903	-09-852-9030 -09-852-9030 -09-852-9030 -09-971-353 -09-971-353 -09-852-9030 -09-852-9030 -10-085-906 -10-085-906 -10-085-906	-10-661-355-24 -10-661-099-24 -10-661-415-24 -10-661-402-24 -10-661-403-24 -10-661-403-24
Sequence 15, Appl Sequence 34, Appl Sequence 29, Appl Sequence 35, Appl Sequence 36, Appl Sequence 450, Appl Sequence 450, Appl Sequence 450, Appl Sequence 2662, Appl Sequence 2662, Appl Sequence 2662, Appl Sequence 2681, Appl Sequence 2881, Appl Sequence 25942, Appl Sequence 25942	equence 25, equence 25, equence 136 equence 31, equence 32, equence 33, equence 12,	nce 26 nce 27 nce 28 nce 33 nce 571 nce 29 nce 28 nce 11 nce 11 nce 36 nce 36	equence 24, equence 24, equence 24, equence 24, equence 24, equence 24, quence 495,

ALIGNMENTS

Sequence 23, Application US/09852903C Publication No. US20030104376A1 GENERAL INFORMATION:

APPLICANT: Diatech Pty. Ltd.
TITLE OF INVENTION: An assay
FILE REFERENCE: 2414918/EJH
CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR FILING DATE: 2000-05-09

US-09-852-903C-23

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PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.0
SEQ ID NO 23
LENGTH: 36
TYPE: DNA
ORGANISM: artificial sequence
FEATURE:
NAME/KEY: misc feature
LOCATION: ()...()
LOCATION: ()...()
                                                                           ; OTHER INFORMATION: CA-19 US-09-852-903C-23
Query Match
Best Local Similarity
Matches 36; Conserv
100.0%; Score 36; DB 10; ilarity 100.0%; Pred. No. 7.1e-05; Conservative 0; Mismatches 0;
                                     Length 36;
   Indels
   °.
 Gaps
 0;
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CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR FILING DATE: 2000-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.0
SEQ ID NO 24
LENGTH: 38
TYPE: DNA
ORGANISM: artificial sequence
PEATURE:
NAME/KEY: misc feature
LOCATION: ()..()
COTHER INFORMATION: CA-20
US-09-852-903C-24
RESULT 4
US-09-263-959-678/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-909-317-7
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                                                                                                                                                              Query Match
Best Local (
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PRIOR APPLICATION NUMBER: 09/280,181
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Betty P. Tsao (Inventor)
APPLICANT: Rita M. Cantor (Inventor)
APPLICANT: Jerome I. Rotter (Inventor)
TITLE OF INVENTION: Genetic Marker Test for Lupus
FILE REFERENCE: 18810-82152
CURRENT APPLICATION UNDERS: US/09/909,317
CURRENT FILING DATE: 2001-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 2414918/EJH
CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: An assay
                                                                                                                                                                Local Similarity
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                                                                                                         1 CACACACACACACACACACACACACACACACACA
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To. US20030104376A1
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                                                                                                                                                              100.0%; Score 36; DB 10; 100.0%; Pred. No. 7.2e-05;
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                                                                                                                                               Mismatches
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Sequence 25, Application US/09852903C
Publication No. US20030104376A1
GENERAL INFORMATION:
APPLICANT: Diatech Pty. Ltd.
TITLE OF INVENTION: An assay
FILE REFERENCE: 2414918/EJH
CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR FILING DATE: 2000-05-09
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR PRIOR DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.0
; FEATURE:
, NAME/KEY: misc feature
; LOCATION: ()...()
; OTHER INFORMATION: CA-21
US-09-852-903C-25
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 39 base pairs
TYPE: nucleic acid
                                                                                                      SEQ ID NO 25
LENGTH: 40
TYPE: DNA
ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIOPPY disk
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V6
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
CLASSIFICATION:
ATTORNEY/AGRAM
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: McMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Rowen, Lee
APPLICANT: KOOD, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
NUMBER OF SEQUENCES: 1279
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and Berry LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (206)
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Length 40;

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US-10-661-097-24
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CURRENT APPLICATION NUMBER: US/10/661,097
CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/1B03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-09-13
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CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 36
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SEQ ID NO 24
LENGTH: 40
                                                                                                       NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATENTIN VET. 3.2
SEQ ID NO 24
LENGTH: 40
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Best Local Similarity
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APPLICANT: JUTEAU, JEAN-MA
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TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES TARGETING HSV
FILE REFERENCE: 029849/0206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: VAILLANT, ANDREW APPLICANT: JUTEAU, JEAN-MAI
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: JUTEAU, JEAN-MARC
OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: oligonuclectide
                                                             ORGANISM: Artificial Sequence
                                                                                    TYPE: DNA
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ORGANISM: Artificial Sequence
                                                FEATURE:
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US-10-661-099-24
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Publication No. US20040171568A1
GENERAL INFORMATION:
APPLICANT: VAILLANT, ANDREW
APPLICANT: JUTEAU, JEAN-MARC
                                                                                               SOFTWARE: PA
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SEQ ID NO 24
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Best Local (
                                                                                                                                                                                                                                 APPLICANT: JUTEAU, JEAN MARC
TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES TARGETING
FILE REFERENCE: 0.29449/0203
CURRENT APPLICATION NUMBER: US/10/661,099
CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
                                                                                                                                    PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 36
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PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 36
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CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR FILING DATE: 2002-12-05
PRIOR FILING DATE: 2002-12-05
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TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES
                        TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                      FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                                                              ENGTH:
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                                                                                                                   PatentIn Ver. 3.2
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No. US20040170959A1
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                                                                                                CURRENT APPLICATION NUMBER: US/10/661,402
CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 36
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                                   SOFTWARE: Patentin Ver. 3.2
SEQ ID NO 24
LENGTH: 40
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SEQ ID NO 24
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: 60/430,934
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-09-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: JUTEAU, JEAN-MARC
TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES TARGETING RSV
FILE REFERENCE: 029849/0205
CURRENT APPLICATION NUMBER: US/10/661,415
CURRENT FILING DATE: 2003-09-12
CURRENT FILING DATE: 2003-09-12
                                                                                                                                                                                                                                                                                               APPLICANT: JUTEAU, JEAN-MARC
TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES TARGETING VIRAL FAMILIES
FILE REFERENCE: 029849/0207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: VAILLANT, ANDREW APPLICANT: JUTEAU, JEAN-MAI
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: VAILLANT, ANDREW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
ORGANISM: Artificial Sequence
                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH: 40
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Local Similarity 100.0%;
Les 36; Conservative 0;
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Pred. No.
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US-09-263-959-495/c
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PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: 60/410,264
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 3.2
SEQ ID NO 24
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Matches
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Patent No. US20020150891A1
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Publication No.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION: APPLICANT: Hood,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/661,403
CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: PCT/IB03/04573
PRIOR FILING DATE: 2003-09-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: JUTEAU, JEAN-MARC
TITLE OF INVENTION: ANTIVIRAL OLIGONUCLEOTIDES
FILE REFERENCE: 029849/0202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: VAILLANT, ANDREW
APPLICANT: JUTEAU, JEAN-MAI
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: oligonuclectide
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TYPE: DNA
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                 APPLICANT: Rowen, Lee
APPLICANT: Koop, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH
NUMBER OF SEQUENCES: 1279
                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                       COUNTRY:
                                                                                                                                                                                                  STREET:
                                                                                                                                                                                CITY: Seattle
                                                                                                                                                                                                                     ADDRESSEE:
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                                                                                                                                                           Washington
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                                                                                                                                                                                                  6300 Columbia Center,
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                                                                                                                                                                                                                       Seed and Berry LLP
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                        Version #1.25
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CLASSIFICATION

APPLICATION NUMBER: US/09/263,959 FILING DATE: 05-MAR-1999

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Sequence 26, Application US/09852903C
Publication No. US20030104376A1
GENERAL INFORMATION:
APPLICANT: Diatech Pty. Ltd.
TITLE OF INVENTION: An assay
FILE REFERENCE: 2414918/EJH
CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR APPLICATION NUMBER: US 60/202,771
PRIOR APPLICATION NUMBER: US 60/202,759
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.0
SEQ ID NO 26
LENGTH: 42
TYPE: DNA
ORGANISM: artificial sequence
FEATURE:
NAME/KEY: misc feature
LOCATION: ()...()
COTHER INFORMATION: CA-22
US-09-852-0900-26
Sequence 27, Application US/09852903C
Publication No. US20030104376A1
GENERAL INFORMATION:
APPLICANT: Distech Pty. Ltd.
TITLE OF INVENTION: An assay
FILE REFERENCE: 24149182H
CURRENT APPLICATION NUMBER: US/09/852,903C
CURRENT FILLNG DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 60/202,771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: MCMSSters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 495:
SEQUENCE CHARACTERISTICS:
LENGTH: 41 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TODOLOGY: linear
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US-09-852-903C-27
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US-09-852-903C-26
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l Similarity 100.0%; Pred. No. 7.2e-05;
36; Conservative 0; Mismatches 0;
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; LENGTH: 44
; TYPE: DNA
; ORGANISM: artificial sequence
; PEATURE:
; NAME/KEY: misc feature
; LOCATION: ()..T)
; OTHER INFORMATION: CA-23
US-09-852-903C-27
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                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2000-05-09
PRIOR APPLICATION NUMBER: US 60/202,559
PRIOR FILING DATE: 2000-05-10
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.0
SEQ ID NO 27
                                                                Matches
                                                                              Query Match
Best Local
                                                              1 Similarity 36; Conserv
               1 CACACACACACACACACACACACACACACACACA 36
CACACACACACACACACACACACACACACACACA
                                                              Conservative
                                                                           100.0%; Score 36; DB 10; 100.0%; Pred. No. 7.2e-05;
                                                                Mismatches
                                                                                              Length 44;
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Search completed: September 2, 2005, 03:54:02 Job time : 39.0082 secs

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Minimum DB
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Perfect score:
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Match
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
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US-09-517-467B-3
US-08-044-618-5
US-08-056-248D-24
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US-09-994-016-14854
US-09-949-016-13423
US-09-949-016-15078
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Compugen Ltd.
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   1, Appli
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17854, A
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16098, A
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11890, A
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618-5 5, Applicat: 10. 5449605 INFORMATION CANT: SMULSO OF INVENTION OF INVENTION OF INVENTION OF INVENTION	0 4 5 4 5 4 5	467B-3 3, Application 6451602 INFORMATION: NT: Ian Popol NT: Lex M. (FERNICE: RTS: FERENCE: RTS: APPLICATION DATE: APPLICATION DATE: APPLICATION NOT SEQ ID NOS OF SEQ ID	222222222222222222
-5 Application 5449605 50000000000000000000000000000000000	larity Conservat TTTCTAGGT TTTCTAGGT TTTCTAGGT GTCTTCGGA GTCTTCGGA	Applicati Applicati Applicati Applicati MetaTion: Ian Popof Lex M. C TOVENTION. LICATION LICA	
N: N: N: N:	8.4 Y 100.1 Ervative PAGGTCGTG AGGTCGTG AGGTCGTG AGGTCGTG AGGTAAG AGGTAAG AGGTAAGAG AGGAGAAGAG AGGCAAGAGC AGCCAAGAGAGC AGCCAAGAGAGC AGCCAAGAGAGC	467B-3 3, Application US/0951746 3, Application US/0951746 20. 6451602 INFORMATION: TI IAN POPOFF WT: IAN COWSETT WT: LEX M. COWSETT WT: LEX M. COWSETT WT: LOY M. COWSET WT: LOY M. COWSETT WT: LOY M. COW	601 2252 2273 8907 113042 152132 152132 152145 177251 16738 16738 16738 16738 40493 41863
US/080446 MARK METHOD OR CANCER BY POLYMORPH	1.4*; 0.0*; 0.0*; 0.00000000000000000000000	/0951 : : ::NSE :: US :: US :: US :-03/5	444444444444444444444444444444444444444
US/08044618 MARK METHOD OR DETECTING A PREDISPOSITION CANCER BY THE USED OF RESTRICTION FRA POLYMORPHISM OF THE GENE FOR THE HUMA POLYMERASE	Score 175; DB 3; Pred. No. 6e-59; 0; Mismatches 0; TCGGGCTTCCGGAGCTTTGG	7B ULATION OF PARP /517,467B	US-09-949-016-161293 US-09-949-016-19 US-09-949-016-11761 US-09-949-016-112343 US-09-949-016-12343 US-09-949-016-12346 US-09-949-016-12371 US-09-949-016-12371 US-09-949-016-12381 US-09-949-016-12168 US-09-949-016-124948
DSITION TO FION FRAGMENT LENGTH THE HUMAN POLY (ADP-RIBOSE)	Length 3660; Indels 0; Gaps 0; CGGCAGCTAGGGGAGGATGGC 1947 CGGCAGCTAGGGGAGGATGGC 164 CGGCAGCTAGGGGAGGATGGC 164 GAGCGGGCGCCCCCTCTTGCAA 2007	EXPRESSION	Sequence 161293, Sequence 4519, Appl Sequence 19, Appl Sequence 11761, A Sequence 16246, A Sequence 15246, A Sequence 1341, A Sequence 1341, A Sequence 13947, A Sequence 199479, Sequence 19168, A Sequence 16168, A Sequence 14168, A Sequence 14578, A Sequence 14578, A Sequence 14578, A Sequence 14578, A Sequence 15527, A Sequence 15527, A Sequence 15453, A Sequence 15453, A

NUMBER OF SEQUENCES: 7

ADDRESSEE:

E: Sterne, Kessler, Goldstein & Fox 1225 Connecticut Suite 300

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SEQ ID NO 24
LENGTH: 3045
TYPE: DNA
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Best Local Similarity
Matches 175; Conserv
                                                                                                                                                                                                                                                                                                                                                      Sequence 24, Application US/09596248D Patent No. 6599727
                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                           APPLICANT: Christenson, Erik
APPLICANT: DeMaggio, Anthony J
APPLICANT: DeMaggio, Anthony J
APPLICANT: Goldman, Phyllis S
APPLICANT: McElligott, David L
TITLE OF INVENTION: Human Poly(ADP-Ribose) Polymerase 2 Materials
FILE REFERENCE: 27866/36544
CURRENT APPLICATION NUMBER: US/09/596,248D
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: 60/139,543
FRIOR FILING DATE: 1999-06-16
NUMBER OF SEQ ID NOS: 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/044,618
FILING DATE: 1930406
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/257,696
FILING DATE: 14-OCT-1988
ATTORNEY/AGENT INFORMATION:
NAME: FOX, SAMUEL L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (202)833-8716
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                              SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 30,353
REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)466-0800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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STATE: L.
20036
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TYPE: NUCLEIC ACID
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1948 GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCCCCTCTTGCAA
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100.0%; Pr
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; Pred. No. 6e-59;
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US-08-860-886-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Applic
Patent No. 6335009
Query Match
Best Local Similarity
Matches 173; Conserv
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Best Local Similarity
                                                                                                                                                                                                          TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 3792 base pair
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NAME/KEY: CDS
LOCATION: (1).
                                                                                                      FEATURE:
NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/860,886
FILING DATE: 03-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Burkle, Alexander
APPLICANT: Zur Hausen, Harald
APPLICANT: Jan-Heiner, Kupper
TITLE OF INVENTION: VECTORS AND VIRUSES FOR USE
TITLE OF INVENTION: IN GENE THERAPY
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                TELEPHONE: 650-493-4935
                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
                                                                                       NAME/KEY: Coding Sequence LOCATION: 96...3134
OTHER INFORMATION:
                                                                                                                                                                          LENGTH: 3792 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 84
                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE:
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                                                                                                                                                                                                                                                                                TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2003 TGCAAGAAATGCAGCGAGAGCATCCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Pennie & Edmonds, LLP
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                  650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                             linear
                   3.5%;
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 Score 73; DB 3;
Pred. No. 3.1e-19;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                    8484-0028-999
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                                   Length 3792;
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   Gaps
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FILE REFERENCE: 27866/36544

CURRENT APPLICATION NUMBER: US/09/596,248D

CURRENT FILING DATE: 2000-06-16

PRIOR APPLICATION NUMBER: 60/139,543

PRIOR FILING DATE: 1999-06-16

NUMBER OF SEQ ID NOS: 68

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 46

LENGTH: 3200
                                                                                                                                                                                                                                                                                                                                                   RESULT 6
US-08-044-618-7
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US-09-596-248D-46
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                                                                                                                                                                                                                                                                                                           Sequence 7, Application US/08044618 Patent No. 5449605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 3.4%;
Best Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Christenson, Erik
APPLICANT: DeMaggio, Anthony J
APPLICANT: Goldman, Phyllis S
APPLICANT: McElligott, David L
TITLE OF INVENTION: Human Poly(ADP-Ribose) Polymerase 2 Materials and
TITLE OF INVENTION: Methods
                                                                                                                                                                                                                                                                     GENERAL INFORMATION: APPLICANT: SMULSO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:hPARP1/hPARP2
OTHER INFORMATION: Fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                               APPLICANT: SMULSON, MARK
TITLE OF INVENTION: METHOD OR DETECTING A PREDISPOSITION TO
TITLE OF INVENTION: CANCER BY THE USED OF RESTRICTION FRAGMENT LENGTH
TITLE OF INVENTION: POLYMORPHISM OF THE GENE FOR THE HUMAN POLY (ADP-ITITLE OF INVENTION: POLYMERASE
                           CITY: TO C. C. STATE: D.C. 20036
                                              STREET: 1225
STREET: Mashington
CITY: Washington
TMATE: D.C.
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                           NUMBER OF SEQUENCES: 7
                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             2003 TGCAAGAAATG 2013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1943 ATGGCGGAGTCTTCGGATAAGCTCTATCGAGTACGCCAAGAGAGCGGGCGCCCCTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1948 GGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCCGGGCGCGCCTCTTGCAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATGGCGGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCGCCTCT 168
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                                                                                                     E: Sterne, Kessler, Goldstein & 1225 Connecticut Suite 300
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOUTWARE: FASTSEQ FOR WINDOWS Version 4.0
SEQ ID NO 14854
LENGTH: 8848
TYPE: DNA
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                                                                               Query Match
Best Local S
Matches 50
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (202)833-871
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: CL001307
                                                                                                                                                                                 ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/257,696
FILING DATE: 14-OCT-1988
ATTORNEY/AGENT INFORMATION:
NAME: FOX, SAMUEL L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 30,353
REFERENCE/DOCKET NUMBER: 0654.0490001
TELECOMMUNICATION INFORMATION:
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TYPE: NUCLBIC ACID
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (202)466-0800
TELEFAX: (202)833-8716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 199304 CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1998 CCTCTTGCAAGAAATGCAGCGAGAGCATCCCCCAAGGACTCGCTCCGGATGGCCATCATGG
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4278 GGTGGGAGGATTGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACAT 4229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        828 CCTCTTGCAAGAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGG
                                    770 GGTGGGAGGATTGCTTGAGGCCAGGGGTTCAAGACCAGCCTGGGCAACAT 819
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                                                                               Conservative
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E: DNA
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                                                                                             2.4%; Score 50; DB 4; L
100.0%; Pred. No. 2.6e-10;
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100.0%; Pred. No. 9.6e-16;
n. Mismatches 0;
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                                                                               Mismatches
                                                                                                                   Length 8848;
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                                                                               Gaps
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04.14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING
RESULT 10
US-09-949-016-13423/c
; Sequence 13423, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Human US-09-949-016-178058
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US-09-949-016-178058
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR PILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FRAESEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 178058
LENGTH: 601
                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local
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LENGTH: 601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                         46;
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Similarity 100.0%; F
46; Conservative 0;
                                                                                                                                                                                                                                                                       GGTTTTGCCATGTTGTCCAGGCTGGTCTTGAACTCCTGGGCTCAAG 347
                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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100.0%; Pr/
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Pred. No. 1.3e-0
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Pred. No.
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; ORGANISM: Human
US-09-949-016-13423
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES AS:
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF
TITLE REFERENCE: CLU01307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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                                                                                                                                                ; Sequence 15078, Application US/09949016
; Patent No. 6812339
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                    GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
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SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16847
LENGTH: 98302
TYPE: DNA
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
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SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 13423
LENGTH: 69909
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Best Local Similarity
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Best Local Similarity
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
CURRENT APPLICATION NUMBER: US/09/949,016
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100.0%; Pred. No.
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100.0%; Pred. No.
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7.3e-09
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OF DETECTION AND USES THEREOF
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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Dioclert, A.
APPLICANT: Glordano, J.Y.
ITILE OF INVENTION: Expressed Sequence Tags and Enc.
Patent No. 6783961
PILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1399-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
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CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 16998
LENGTH: 73
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US-09-513-999C-16098
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Best Local Similarity
Matches 46; Conserva
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US-09-513-999C-16110
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US-09-949-016-15078
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PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                Sequence 16110, Application US/09513999C Patent No. 6783961 GENERAL INFORMATION:
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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and
Patent No. 6783961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 15078
LENGTH: 114426
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Local Similarity 100.0%; Pred. No. 3.9e-08;
les 45; Conservative 0; Mismatches 0;
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Pred. No. 6.9e-09;
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Search completed: September Job time : 381 secs

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APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags
Patent No. 6783961
FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 29990
                                                    Matches
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Best Local Similarity
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LENGTH: 74
                                                                  Query Match
Best Local
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CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature LOCATION: 16
                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                     NAME/KEY: misc_feature LOCATION: 86
                                                                                                                                                                FEATURE:
                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: 18
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OTHER INFORMATION: 8=9
                                                                                                                      OTHER INFORMATION: k=g
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                                                                  Local Similarity
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                        1398 GCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGTGGGAGGATCGCT 1442
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100.0%; Pred. No.
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